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Date: 07/27/99

LARGE-SCALE CsCl preparation of Bacterial genomic DNA

1. Grow 100 ml culture of bacterial strain to saturation.
2. Pellet cells for 10 min at 6000 rpm (use SSA rotor)
3. Resuspend cells gently in 9.5 ml TE buffer. Add 0.5 ml of 10% SDS & 50 μ l of 20 mg/ml proteinase K. Mix thoroughly and incubate 1 hr at 37°C.
4. Add 1.8 ml of 5M NaCl and mix thoroughly.
5. Add 1.5 ml CTAB/NaCl solution. Mix thoroughly and incubate 20 min at 65°C. (can stay for a moment)
6. Add an equal volume (13 ml) of chloroform/isoamyl alcohol. Extract thoroughly. Spin 10 min at 7000 rpm (use SS34, room temperature) to separate phases.
7. Transfer aqueous supernatant to a fresh tube using a wide-bore pipet.
8. Add an equal volume phenol/chloroform/isoamyl alcohol. Extraction the equal volume. Repeat three times.
9. Add an equal volume of chloroform/isoamyl alcohol.
10. Add 0.6 volume isopropanol and mix. (-20°C overnight) (-20°C can help to precipitate.)

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2

Continue:

11. Transfer the precipitate to ~~1~~ 1 ml of 70% ethanol in a fresh tube (just brown-red one), by hooking it on the end of a Pasteur pipet that has been bent (by heating it) and sealed.
12. Drying the pellet (just in the air). Remove ~~start~~ white and stringy DNA to the 4th 2nd TE buffer.
13. Do the electrophoresis of nucleic acid (there are a lot of DNA and RNA)
14. Use RNase to digest the genomic DNA overnight.

Genomic DNA. (continue)

1. Use *EcoR* I to do restriction. put

{	genomic DNA	11 μ l
	buffer (1x)	9 μ l
	<i>EcoR</i> I	4 μ l

 overnight.

Digest genomic DNA.

2. do the electrophoresis at 60 V.

put 109 μ l. digested DNA (90 μ l. digested DNA
19 μ l dye)

the result is not good.

Extracting Plasmid DNA of *E. coli*. (Qiagen Midi and Maxi Prot.)

1. Inoculate a 5-ml LB medium with *E. coli*. 37°C. shake. to afternoon.
2. Inoculate 200ml medium. Grow at 37°C ~~for~~ overnight.

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Extracting plasmid DNA. (continue)

3. Harvest the bacterial cells by centrifugation at 6000g for 15 min at 4°C.
4. Resuspend the bacterial pellet in 4 ml of buffer P1.
5. Add 4 ml of buffer P2, mix gently but thoroughly by inverting 4-6 times, and incubate at room temperature for 5 min.
6. Add 4 ml of chilled buffer P3, mix immediately but gently by inverting 4-6 times, and incubate on ice for 15 min.
7. Centrifuge at 20000 g for 30 min at 4°C. Remove supernatant containing plasmid DNA promptly.
8. Re-centrifuge the supernatant at 20,000 xg for 15 min at 4°C. Remove supernatant containing plasmid DNA promptly. ~~Alternatively,~~
~~the sample can be fill~~
9. Equilibrate a QIAGEN-tip 100 by applying 4 ml Buffer QBT, and allow the column to empty by gravity flow.
10. Apply the supernatant from step 8 to the QIAGEN-tip and allow it to enter the resin by gravity flow.
11. Wash the QIAGEN-tip with 10 ml (two times) buffer QC.
12. Elute DNA with 5 ml buffer QF.
13. precipitate DNA by adding 3.5 ml room-temperature isopropanol to the eluted DNA. Mix and Centrifuge immediately at ^{13000 rpm} ~~15000 g~~ for 30 min at 4°C. Carefully decant the supernatant.
14. Wash DNA pellet with 2 ml of room-temperature 70% ethanol, and centrifuge at ^{13000 rpm} ~~15000 g~~ for 10 min. Carefully decant the supernatant without

disturbing the ~~pette~~ pellet.

15. Air-dry the pellet for 5-10 min. and redissolve the DNA in a suitable volume of buffer \rightarrow (TE buffer \rightarrow 1/2 ml).

16. Do electrophoresis to check it. \rightarrow to digest? \rightarrow next page

8/3/99

High-efficiency transformation by electroporation

1. Inoculate a single colony of E. coli cells into 5 ml LB medium. Grow 5 hr. to overnight at 37°C with moderate shaking.

2. Inoculate 2 ml of the culture into 50 ml LB medium in a sterile flask. Grow at 37°C shaking at ^{shaker} in incubator to an OD₆₀₀ of ~~0.5-0.7~~ 1.25. (if > 0.7 need more time)

3. Chill cells in an ice-water bath 10 to 15 min and transfer to a prechilled 50 ml centrifuge bottle.

4. Centrifuge cells ^{10 min} ~~20 min~~ at ⁸⁰⁰⁰ ~~6000~~ rpm. 2°C.

5. Pour off supernatant and ~~resuspend~~ resuspend the pellet in ice-cold water. Add 25 ml ice-cold water and mix well.

Centrifuge cells as in step 4. ^{at 10 min 8000 rpm. repeat 2 times.} (throw away the supernatant to the bleach ^{bottle including})

6. Pour off supernatant immediately and resuspend the pellet by swirling in remaining liquid.

7. Add another ml ice-cold water, mix well, and centrifuge again in step 4.

8. Pour off supernatant immediately and resuspend the pellet by swirling in remaining liquid.

9. Put remaining liquid to the eppendorf tube. centrifuge it at 12000 rpm 5 min. pour off by using vacuum.

10. add 100 ul water to the tube. mix well. put it on ice.

(合併管カ管)

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4

6

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- x10. Set the electroporation apparatus to 2.5 kV, 25 μ f. Set the pulse controller to 200 or 400 ohms? (Under the machine instruction)
11. Add 1 μ l plasmid DNA (PGP 704) ^{and 20 μ l bacterial} in _____ to tubes containing fresh or thawed cells (on ice). Mix by tapping the tube or by swirling the cells with the pipettor.
12. Transfer the DNA and cells into a uvette that has been chilled 5 min on ice. Shake slightly to settle the cells to the bottom, and wipe the ice ^{between} and water from the uvette with a Kimwipe.
13. Place the uvette into the sample chamber.
14. Apply the pulse by pushing the button of flipping the switch.
15. Remove the uvette. Immediately add ^{LB broth} ~~SOG medium~~ ^{immediately} and transfer ^{it} to a sterile culture tube with a Pasteur pipet. Incubate ~~30~~ to 60 min with moderate shaking ^{37°C}.
16. Plate aliquots of the transformation culture on LB plates containing antibiotics.

plasmid digest100 μ l

08/03/99

PGP 100	1 μ l
10x buffer	1 μ l
DNasefree H ₂ O	7.5 μ l
ECOR I	0.5 μ l

37°C 2 hr. \rightarrow 65°C for 15 minrun 5 μ l on geldilution 10^{-6} 100 μ l to LB DAP 10^0 10^{-2} 100 μ l to LB DAP DAP

result

 10^0 10^{-2} zero 2-3 μ l 10^{-6} 100 colonies

Do electrophoresis.

1. 15 ~~ul~~ marker (1kb) first line
2. 65 ~~ul~~ digested DNA + 13 ~~ul~~ 6x dye. Second line.
3. Cut out 7-10 kb gene.
4. elute into Bio-rad - promega.

Fragment Isolation Protocol from NuSieve Gel.

1. Place the electrophoresed gel on the UV transilluminator and view under long wave light. Locate the fragment of 7-10 kb and cut out the band using a glass cover slip and place the gel fragment into a 1.5ml microcentrifuge tube.
2. Heat to 65°C until gel slice melts.
3. Add ^{120 ul} 3 volumes of BioRad DNA isolation binding buffer and for every 2 ^{1 ul} ug of DNA add 5 ^{30 ul matrix} ul of matrix.
4. Attach a clean 3cc syringe to a promega spin column and add the solution to it.
5. Push this slowly through the column (1 drop/3 sec)
6. Detach column from syringe and remove plunger.
7. Wash with 2 mls of 95% isopropanol.
8. Spin at 12,000 rpm for 20 sec.
9. Dry Column in Hybridization oven for a few minutes ^{65°C, 10 min} at 68°C.
10. Add ^{30-50 ul} 30-50 ul of water or TE. wait 1 min and spin again at 12,000 rpm for 20 sec.
11. Run 10 ^{5 ul} ul of the eluant on a gel to check recovery.

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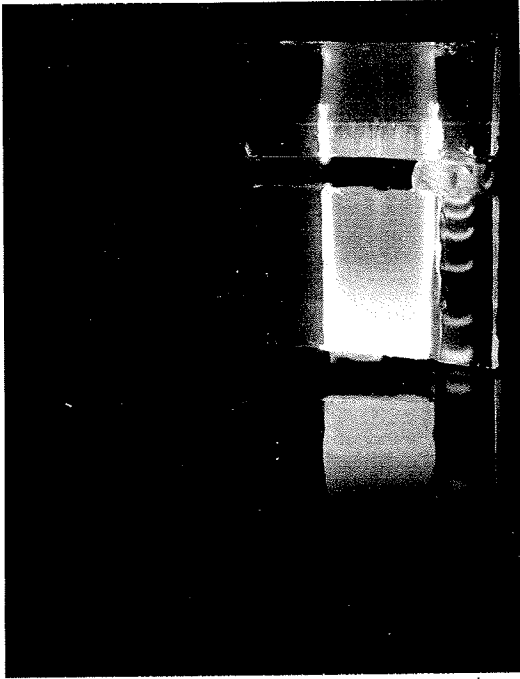
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4

8

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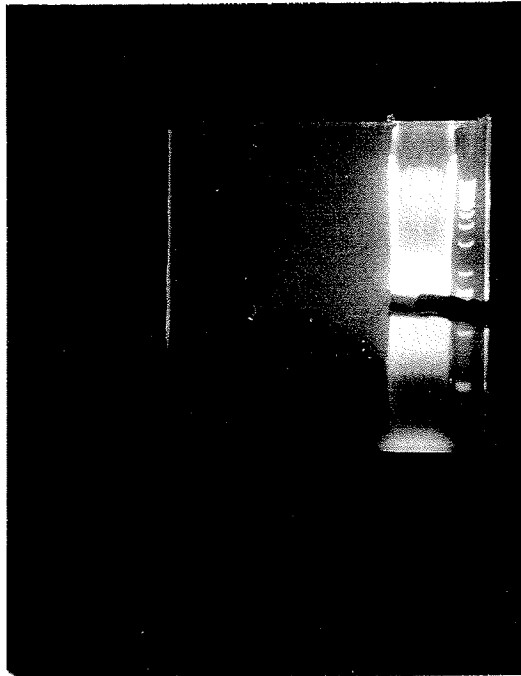


cut out 7-10 kb

6
5
4
3
2

1 → bright 1 kb

digested DNA marker (1 kb)



6
5
4
3
2
1

Do electrophoresis to check the result.

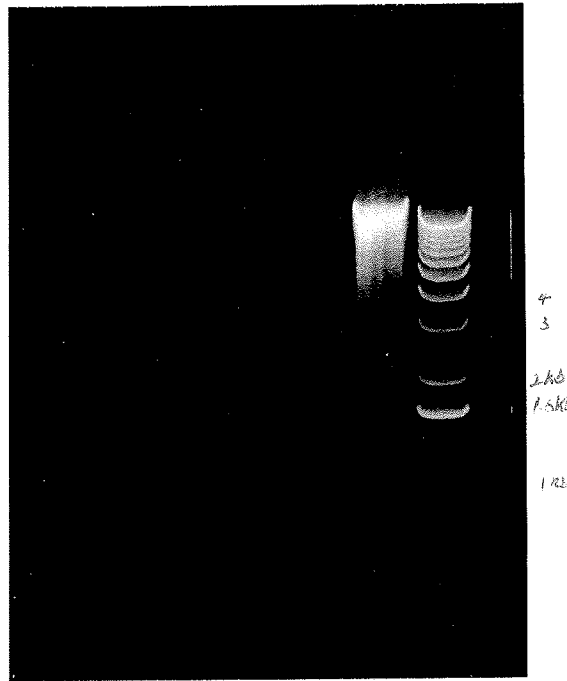
DNA. 5 μ l. + 1 μ l dye.

15 μ l

Marker

\rightarrow 60V. run

Results:



Conjugation:

1. Donor Strain (*E. coli* MGN 617 PGP704 or MGN 617 PLOT-Km) are grown overnight with shaking at 37°C in 2 ml LB broth containing 200 μ g/ml ampicillin and 50 μ g/ml Kanamycin. (DAP)
2. Grown *P. multocida* 11039 overnight with shaking at 37°C.
3. 50 μ l MGN 617 or 50 μ l ~~M~~ + 100 μ l 11039 to 5 ml of 10 mM $MgSO_4$
4. Vortex for a few seconds.
5. Transfer to a 5 ml disposable syringe and filter through a 25 mm Miltrical filter.
6. Drain the filter and carefully remove it from the filter case with sterile forceps (curved tip work best)

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- g. Incubate at 37°C for 8-18 hrs.

2. 36V. Km

08/06/99

~~Results~~ 9. Resuspend the filter using 3ml 10 mM MgSO₄. Vortex.

DGP 704

BHI	Ap	100 μ l	10^0	} negative
BHI	ApK	100 μ l	10^0	
BHI		100 μ l	10^{-5}	
LB	ApDap	100 μ l	10^{-5}	

PLOT-Km

BHI	ApKm	100 μ l	10^0	\rightarrow negative
BHI	Km	100 μ l	10^0	
BHI		100 μ l	10^{-5}	
LB	ApKm	100 μ l	10^{-5}	
		200 μ l		

Transformation by electroporation

PGP 704.

Plot $\text{km} \quad 10^0 \quad 10^{-1} \quad 10^{-2} \quad \text{detention}$

Results

08/09/99

Digestion

93-146 genomic DNA (CL phd/leh... prep)

Leat incubate at 70°C 30min

Restriction Digestion

LECORI

DNA 51 ml

10x buffer 6

F. G. R. I. 3

37°C 4h. → put in 15°C water bath

08/10/99

D. electrophoresis

903-146 genomic DNA (after digest) 60 μ ldye (6x) 10 μ l↓
run gel. (marker 1x6 12 μ l)

Fragment Isolation Protocol from Gel. (See Page 7)

1. Cut out 7-10 kb and place the gel fragment into microcentrifuge tube.
2. Heat to 65°C until gel slice melts.
3. Add 3 volumes (630 μ l) of BioRad DNA isolation binding buffer ~~and buffer~~ (1 μ g \approx 1 μ l volume) and usually add 30 μ l matrix.
4. Attach a clean 3 cc syringe to a promega spin column and add the solution to it.
5. Push this slowly through the column.
6. Detach column from syringe and remove plunger.
7. Wash with 2 mls of 95% isopropanol.
8. Spin at 12,000 rpm for 20 sec.
9. Dry Column in Hybridization oven for a few minutes.
10. Add 30 μ l of TE wait 1 min and spin again at 12,000 rpm for 20 sec.
11. Run Gel.

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29
9130199

Digest:

D1039 60ul

Buffer 7ul

EcoRI 3ul

37°C 2h / 65°C 15 min,

9130199

1. Run a gel. 70ul.
2. Cut out 7-10kb.
3. Use another kit to extract DNA.

QIA quick Gel Extraction kit protocol (another kit)

1. Excise the DNA fragment from the agarose gel with a clean, sharp scalpel.
2. Weigh the gel slice in a tube. Add 3 volumes of Buffer QG to 1 volume of gel (100mg ~ 100ul)
3. Incubate at 50°C for 10 min.
4. After the gel slice has dissolved completely, check that the color of mixture is yellow.
5. Add 1 gel volume of isopropanol to the sample and mix.
6. To bind DNA, pipet the sample onto the QIA quick column and apply vacuum. After the sample has passed through the column, switch off vacuum source.

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Run gel. (Southern Blotting)

1. Dig3 EcoRI Sal I Bam HI Xba I 11039 EcoRI → in back
2. ~~Rinse~~ # ~~Rinse~~
3. photograph with a ruler laid alongside the gel so that band positions can later be identified on the membrane.
3. Rinse the gel in distilled water and place in a clean glass dish containing ~10 gel volumes of 0.25 M HCl. Shake slowly on a platform shaker for ^{10 min} ~~3 min~~ at room temperature.
4. Pour off the HCl and rinse the gel with distilled water. Add ~10 vol denaturation solution and shake as before for 30 min to 1 hr.
5. Pour off the denaturation solution and rinse the gel with distilled water. Add ~10 vol neutralization solution. Shake as before for 30 min to 1 hr.

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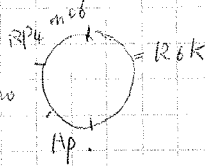
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11/19/2019

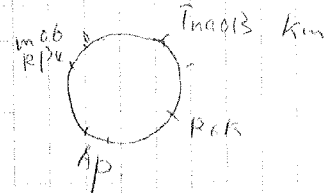
Conjugation:

See P10:

put the conjugation in

① MGN 617 10^{-1} BHI Km② 10^{-1} BHI Ap③ MGN 617 PGP 10^{-1} BHI Ap Km 10^{-1} BHI Km 10^{-1} BHI [Ap] → can grow 10^{-1} BHI Ap Km

③ MGN 617 PLOF Km

 10^{-1} BHI Km x2 (100 μ l. 20-50 colonies) 10^{-1} BHI [Ap] 10^{-1} BHI Ap Km100 μ l 10^{-5} BHI 11039 10^{-5} LB 12ApApKm

→ MGN 617

↓
200 μ l DAP

11/23/99

Cut out 2-6 Kb.

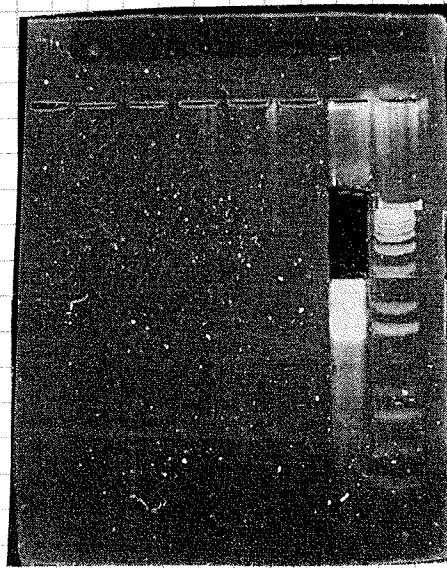
7-10 kb. and more than
10 kb.

use Qiagquick gel

extraction kit to extract

DNA.

result:



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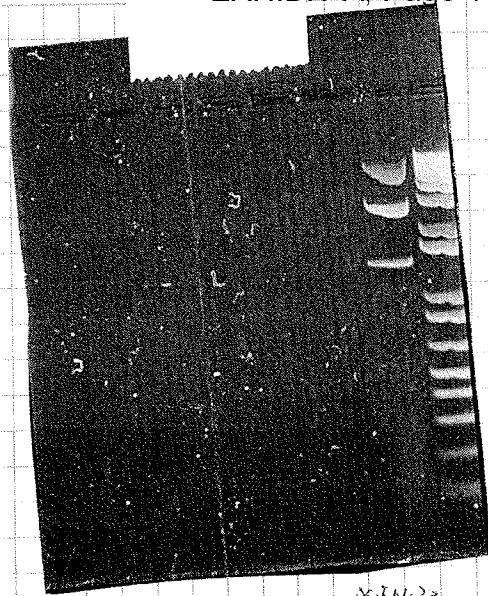
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The digest is not

The pGEM-3Z has not been digested completely.

So there are three bands.
(supercoiled DNA and one strand DNA)



pGEM-3Z
Plasmid DNA (1.1 kb)

So repeat the digest

11039 having 1 week long can not grow in BHI broth.
but can grow in BHI plate.

Inoculate 11039 and HGV 617. PGM 704. PLOT km again.

Do digest again 26 ul DNA + pGEM-3Z)
1 ul EcoRI + 3 ul buffer
over 2h. + 15' 65°C

1113 12000

Sap 1 ul + 4 ul buffer → 37°C 2h. → 15' 65°C

+ 20 ul 100% EtOH + 4 ul 3M NaAc → fraction

+ 70% ~~100%~~ 13500 rpm 15' → pour off supernatant →

100 ul 20% EtOH 13500 rpm 10'. → pour off supernatant
→ dry → run a gel (1 ul)

Conjugation : LB plate + 200 ul OAP + 100 ul IPTG.
do conjugation

BHI → BHI + 100 ul Ap → BHI Ap

BHI + 100 ul km → BHI km

BHI + 100 ul Ap + 100 ul km → BHI Ap km

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2/4/00

~~Run a gel.~~

Digest

21.5 μ l DNA2.5 μ l buffer1 μ l EcoRI

overnight 18/07

 $\rightarrow 65^{\circ}\text{C } 15'$

Run a gel.

cut out

2-6 kb

6-10 kb

more than 10

(because the extraction DNA always bigger than
it is. so, cut out ^{was} smaller band)
little

2/7/00

extract DNA

(use Qiagenick Spin Handl^{kit}er)

2/8/00

Run a gel.

but

no DNA

improve:

Clean DNA

before

to volume 3M Na acetate

2 vol 100% EtOH Clean DNA

 $-20^{\circ}\text{C } \geq 1\text{h}$ centrifuge $\geq 13,000\text{ rpm } 4^{\circ}\text{C } 15'$

Remove supernatant

Add 100 μ l 70% EtOHcentrifuge $\geq 13,000\text{ rpm } 4^{\circ}\text{C } 10\text{ min}$

Remove supernatant

Air dry 5 min

Resuspend 30 μ l DNase free H₂O

Run a gel

2/10/00

★ Clean DNA

 $\frac{1}{10}$ volume 3M Na acetate

2 vol 100% EtOH

 $-20^{\circ}\text{C } \geq 1\text{h}$ centrifuge $\geq 13,000\text{ rpm } 4^{\circ}\text{C } 15'$

Remove supernatant

Add 100 μ l 70% EtOHcentrifuge $\geq 13,000\text{ rpm } 4^{\circ}\text{C } 10\text{ min}$ Remove supernatant. ^{air} dry $\sim 5\text{ min}$ Resuspend 30 μ l DNase free H₂ORun a gel \rightarrow cut out

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3/13/00

1. Run a gel (genomic DNA 11039)

a. Do digest

just has a little DNA
may because the DNA doesn't
dissolve well.

3/14/00

Set up PCR reaction

	①	②	③	④
primer SC 1011	0.8		0.8	0.8
primer SC 1012	0.8		0.8	0.8
dNTP	4	4	0	4
3.3x buffer	6	6	6	6
Mg	2.4	2.4	2.4	2.4
H ₂ O	<u>6</u>	<u>7.6</u>	<u>10</u>	<u>6</u>
	20ul	20ul	20ul	20ul

higher layer

Control DNA	0	0	0	0.5
3.3x buffer	9	9	9	9
enzyme	1	1	1	1
H ₂ O	<u>20</u>	<u>20</u>	<u>20</u>	<u>19.5</u>
	30ul			

put 11039 in 50° for 2 hrs

3/14/00

because a lot of

more than
DNA do not dissolve in TE buffer.

Run a gel (11039 genomic DNA)

① Line 1ul DNA
② Line 3ul DNA

Do digest (11039) 7.5 ul total

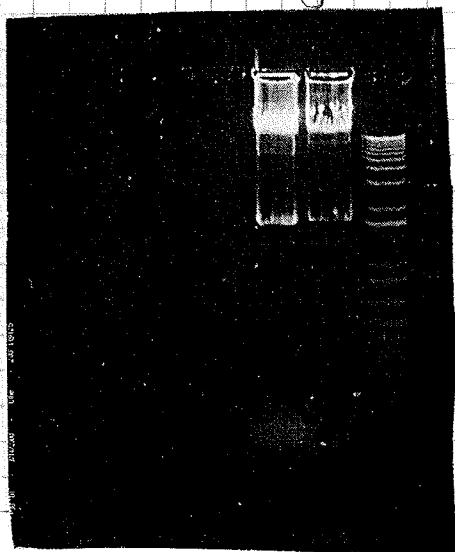
62.5 ul DNA

7.5 ul buffer

5ul Enzyme

37° overnight

65° 15'



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in 3 sq. rubber maid dishes put 2 pcs chromatograph filter paper (this is too soak up buffers)

- 1) Denaturation solution - takes very little just enough to soak up filter. 3 min.
- 2) Neutralization buffer 5 min.
- 3) 2x SSC 5 min. place it on the paper and dry for 30' cross link on program C3 → put them in plastic.

4113100.

1. preheat the hybridization buffer to 42°C.
2. In a suitable container prewet the blot in 5xSSC. Loosely roll the blot and place inside the tube. Add ^asmall amount of 5xSSC to the tube and "unroll" the blot ensuring no air bubbles are trapped between the membrane and the tube. Don't allow the blot to overlap itself.
3. Pour off the 5xSSC and add the appropriate volume of ~~the~~ hybridization (200ul). buffer.
4. Prehybridize in oven for at least 30 min at 42°C.
5. prepare the labelled nucleic acid probe as instruction P13.
 - ⓐ Dilute the DNA to be labelled to a concentration of 100ug/ml using the water supplied.
 - ⓑ Denature 100ug of the DNA sample (100ul) by heating for 5 min in a boiling water bath. ^{try 2ul + 8 gene} (3ul dam + 7ul H₂O)
 - ⓒ Immediately cool the DNA on ice for 5 min. ^{1-10-2 (dam gene)} Spin briefly in a microcentrifuge to collect the contents at the bottom of the tube.
 - ⓓ Add an equivalent volume of DNA labelling reagent (100ul) to the cooled DNA. Mix gently but thoroughly.

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5/14/00

prepare sequencing for genomic DNA

Set up sequence program. (Liang sequence)

Roe-Lab	Big-dye	30-30mer primer	purified gen DNA	H ₂ O
1x reaction	8ul	(10moles)	11039 base	5ul
		1ul (30pmol/ul)		

total 20ul

p.s. primer. (add 3ul primer to 97ul H₂O)

HAPDAM P2

HAPDAM M2

↓
PCR

5/2/00 prepare sequencing

1. Column Hydration

- ① Remove the top of column, then add 0.8ml of reagent grade water. ^{invert for a few times. make sure no bubbles} leave the column for at least 2 hrs. at room temp.

② Removal of Interstitial Fluid:

- ① remove the top cap first, then remove the end stopper from bottom.
- ② allow excess column fluid to drain into a wash tube. discard this fluid.
- ③ spin the column and wash tube in centrifuge at 750g for 2 mins. discard it.

3. Sample processing:

- ① Add 20ul ^{sequence(PCR)} ~~PCR~~ production into tube column. ^(it's on another tube) make sure it's be into the column, but don't touch the column.
- ② spin the column at 750g for 2 min. the purified sample will collect in the collection tube.
- ③ Dry the sample in a vacuum centrifuge.

4. ~~to~~ resuspend sample in 25ul of template reagent in ..

- ① vortex 30sec } X2 2min denature in 95°C
quick spin } chill 5min

put 10ul in sequencing tube

freeze & resuspend 15ul at -20°C

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H.O. sequencing didn't work.

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$17800 \frac{\text{col}}{100} \Rightarrow 178000 \text{ colonies / ml}$

$$\frac{1.78 \times 10^5}{0.005 \text{ ug}} = 3.56 \times 10^7 \text{ / ug DNA}$$

Result: the problem is xgal. the old xgal is not useful for producing blue colonies.

Do electroporation again. and inoculate the restreaked bacteria to the LB Broth.

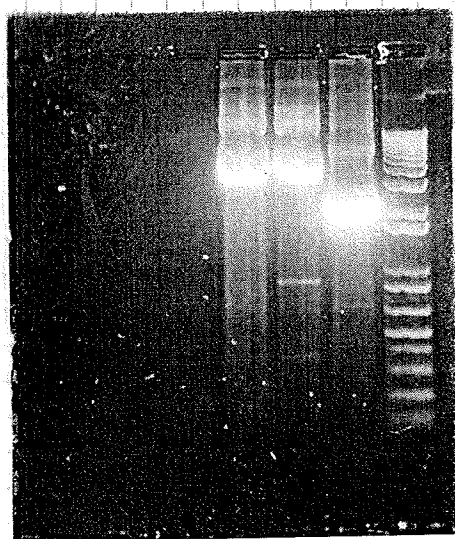
6113/00

Extract the plasmid DNA (restreaked Bact.)

put the electroporation to refrigerator

Run a gel (plasmid DNA ~~EE~~ ① ② ③)

2ul + 3ul + 1ul dye
DNA H₂O



③ ② ①

② and ③ is definitely inserted. due the digest.

20 ul. total

1 ul DNA

2 ul buffer

0.5 ELOR I

16.5 ul H₂O

overnight 37°C load 3-4 ul on a gel.

6114/00

Do ~~electroporation~~ southern hybridization:

Run a gel (plasmid DNA ELOR I digested (20 ul))

load 3-4 ul per well.

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Date: _____

★ 2x SSC 5 min x2 at room temp.
 5x SSC 0.5% SDS 42°C 30' 转 oven
 50 ml 2.0 ml 10%

2x SSC 用报纸 (Kimwipes) 将膜在石板上擦。
 3x SSC 洗干净 rinse and shake 5' ^{4 min} x 2 ~~10'~~

7/19/00

Do ligation

1 μ l PGEM-32 EcoRI + Ssp +rt.
 8 μ l insertor (11039 3-5kb extraction)
 1 μ l buffer
 0.5 μ l ligase (New England Lab)
 16°C overnight

7/10/00

65°C 15'

Do electroporation.

Note: cuvette should be chilled 5 min on ice.
 immediately add 1 ml LB medium and transfer
 to a sterile culture tube.

7/28/00

Run a longer gel for 11039 EColI.

1 lane: ladder (2ul) | 5:30 pm. 20V.

3 lanes: 11039 EColI.

5 ---: ladder (4ul)

It goes very slowly. I should run this gel at 30V.

12:45 end.

1:10 continue

6:10 en

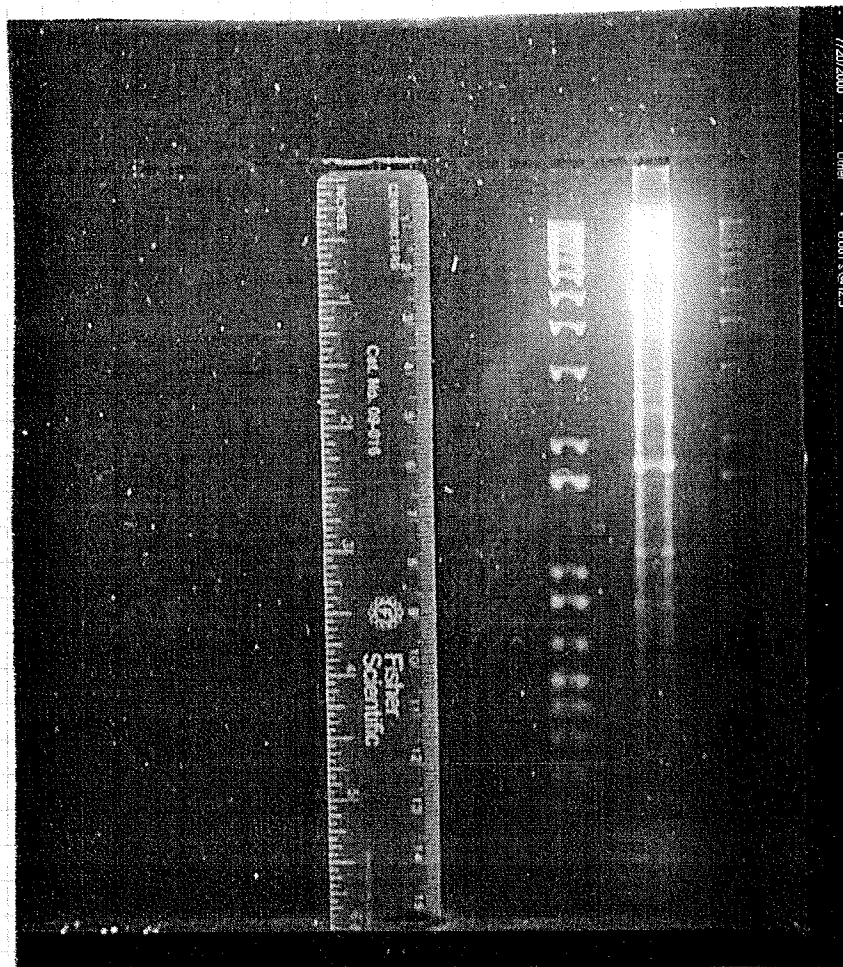
gel size 6x15

paper towel 8x15

filter paper 3 (⁵⁵5x14)

filter paper 5 (8x15)

membrane 7x14



the part from 6 kb to 8 kb has been seperated. Do Southern hybridization.

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Make hybridization buffer.

100ml buffer

5 g blotting reagent.

0.5 mol NaCl 2.922 g.

room temp. stir for 1 hr.

then 42°C at least 30'.

8/03/00

Continue Southern hybridization and Sequencing.

8/04/00

Sequencing didn't work again, but Southern hybridization did

work.

Set up the PCR reaction to make more 1-10-2 and check the plasmid DNA.

primer dilution

95 µl H₂O

2.5 µl HAPdamM1

2.5 µl --- PI

100 µl

→ 1 µl

PCR reaction

① 1-10-2 0.5

dNTP 5

primers 1

buffer 5

tag 0.5

H₂O 38.5

② plasmid 1 µl

dNTP 5

primer 1

buffer 5

tag 0.5

37.5

③ 1-8-1 2

dNTP 5

primers 1

buffer 5

tag 0.5

H₂O 36.5

④ Control 0

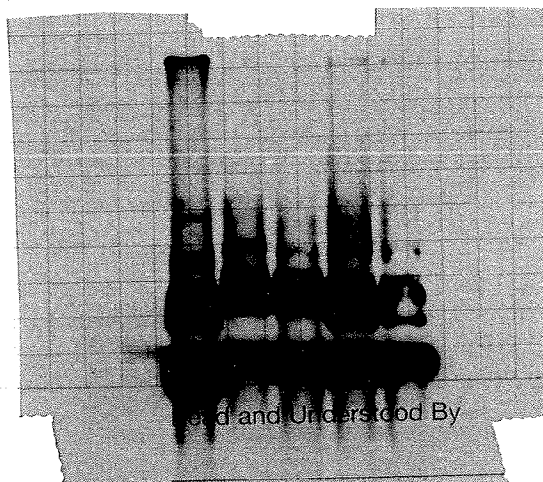
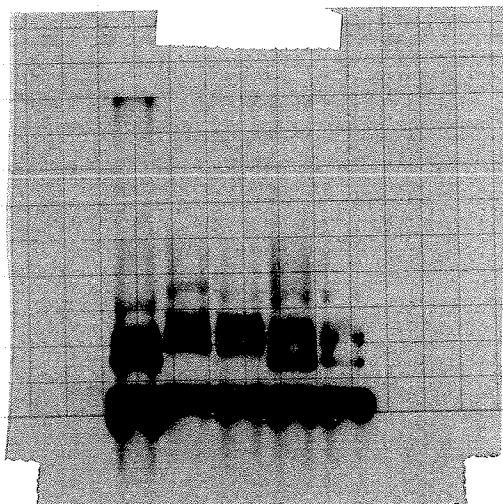
5

1

5

0.5

38.5



Date: _____

8/25/00

Run a sequencing reaction

big dye 8 ul
plasmid 2 ul
primer 2 ul
H₂O 8 ul

Should keep

~~some~~ a little

11039 Hind III

into

21

Left for Southern in

in region

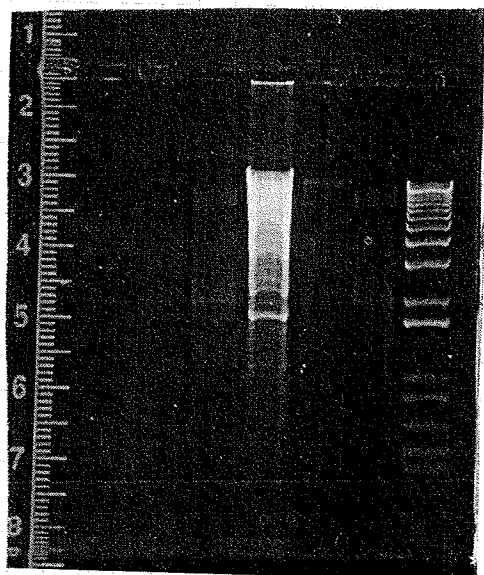
Run a gel for genomic DNA
(25%) in order to ^{comp}sepe
of ladder

8/28/00

Run a gel. 3ul for extracted DNA
3ul for ELOR I 11039 digest (positive control)

- ① extracted DNA 4-6 kb
- ② " " 6-9 kb
- ③ 11039 ELOR I
- ④ repeat ① → another tube
- ⑤ repeat ③ → another tube

8/29/00



the size of both tubes is bigger
than needed. So extract DNA
from 3-4 gel slice.

⑤ ④ ③ ② ①

8/30/00

Run a gel

10/9/00

Digest clone - a - a plasmid DNA to get probe

3ul
Digest 3ul DNA
3ul buffer
1ul ~~ECORI~~ ~~and~~ ~~ECORI~~
23ul H₂O

30ul → 2h. 37°C

15' 65°C

cut out from 1.4 kb.

PGEM-32 HindIII.

10/10/00

1. Continue plasmid DNA + sap treatment
2. do gel extraction.

10/11/00

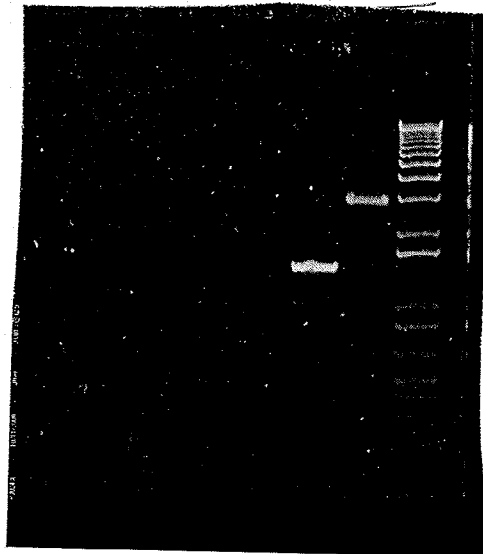
Run a gel for gel extraction (for probe) and
PGEM-32 HindIII + sap trt. (both are 5ul)
gel extraction (probe) just need to use 3ul
for detect.

I use PBS 1.5ul

Insert 7.5ul

buffer 1ul

Ligase 0.5ul



10/12/00

the electroporation didn't work. try
electroporation.

1ul PGEM32 + 10⁻² 10⁻⁴LBAP 10⁻² 10⁻⁴

10/11/00 it did work.

Read and Understood By

Signed

Signed

1/31/01

Do per again

	①	②	③
P.mutacida A13	0	1	1
dNTP	5	5	5
primer	1	1	0.5
buffer	5	5	5
New tag	0.5	0.5	0.5
H ₂ O	38.5	37.5	38.0
50 ul			

95°C 2 min

95°C 30 sec.

72°C 10 min

4°C hold

64°C 1 min

72°C 2 min

1 cycle

35 cycles

2/01/01

Do PCR again. Add
more genomic DNA and
increase
~~decrease~~ annealing
temperature to 66°C

there're
a little DNA on
the gel. It's
not clear. So
Add more genomic
DNA to do PCR
again

	①	②	③	④
P.mut (A13)	0	3	6	10
dNTP	5	5	5	5
primer	1	1	1	1
buffer	5	5	5	5
Tag	0.5	0.5	0.5	0.5
H ₂ O	38.5	35.5	32.5	28.5
50 ul				

digest

EXHIBIT A, Page 27

DNA

buffer

ECORI

H₂O

①

②

③

1

1

1

5

5

5

0.5

0.5

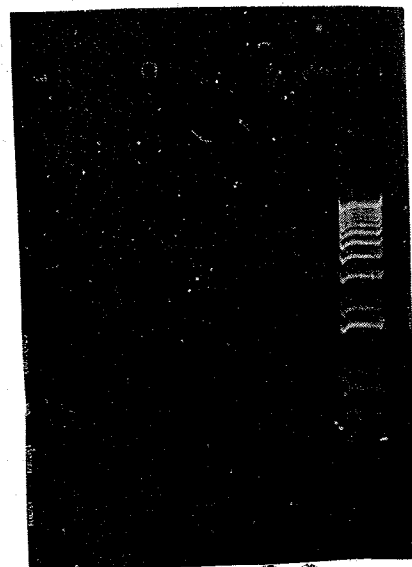
0.5

43

43

43

50 ul



① ② ③

annealing temperature 66°C 1'

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Signed

3/5/01

1. Run a gel for H digest.

2. Do a Klenow for clone A ~~EE~~ clA1 and T-H111
digest

Klenow: 20- μ l digest (the concentration of my DNA is high enough)

0.8 μ l dNTPs (200 μ l of each in working solution)

1 μ l 0.1 mg/ml BSA (I skip this step)

0.5 μ l Klenow

I didn't add buffer because the Multi Core (buffer) is compatible for Klenow

incubate @ 37°C for 30 min. HI @ 75°C for 10 min.

(I lost this Klenow product)

3/6/01

Do digest for B. C. E. F. G. H. and Rnd a gel for it.

I select B. C. G. to run a seq. reaction again.

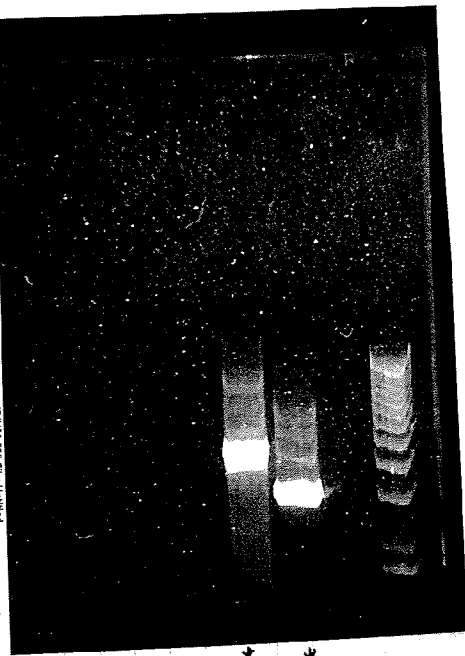
And to make sure which one is right.

Read and Understood By

Signed

Signed

Exp: 10 sec Bin: 11 Gain: 10.00 W5555 60.55 No Date: 6/4/2011 Time: 2:38:33 pm [C:\480-586 File: Untitled



PPS88⁺ MGN617

615101.

Do the digest for PPS88 (Xba I + Kpn I)

5 ul DNA

12 ul H₂O

(multicore) buffer 2 ul

0.5 ul Xba I

0.5 ul Kpn I

2h 37°C

Run a gel for digestion, genomic DNA

PCLPM2

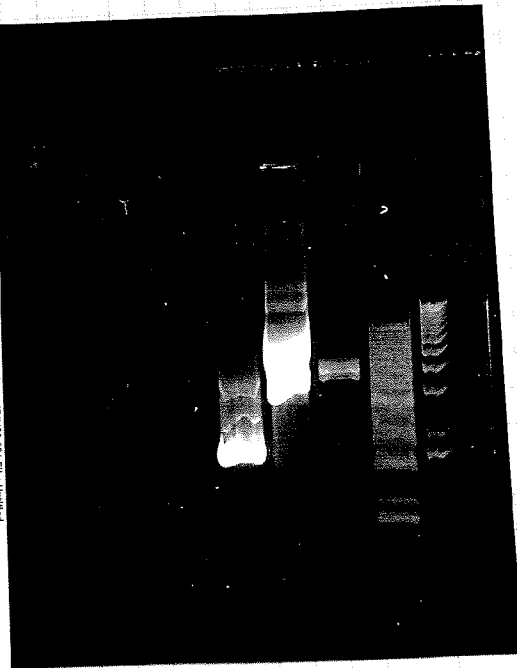
(from clone A)

and pbluescript plasmid

all the samples → 1 ul

inactivate 65°C 15'

Exp: 10 sec Bin: 11 Gain: 10.00 W5555 60.55 No Date: 6/4/2011 Time: 3:04:54 pm [C:\480-586 File: Untitled



PPS 88 clone A MGN.
plasmid

Read and Understood By

Signed

Signed

do sequencing react: for
number 5

Run a gel for ① ③ ⑤

big eye 8 μ l
plasmid 2 μ l

primer T3 2 μ l

water 8 μ l

Sample	Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
bluescript and dam	dsDNA	3.631	5.554	3.175	0	1.74929	0.2777 μ g/ μ L	1	50	174.93

6/29/01

number 5

Sample	Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
PLS 88	dsDNA	0.577	1.176	0.606	0	1.94059	0.0588 μ g/ μ L	1	50	194.06

8 μ l PLS 88

XbaI / BamHI 0.5 μ l

buffer (multicore) 1 μ l

water 0.5 μ l

8 μ l PLS 88

XbaI 0.5 μ l

BamHI 0.5 μ l

multicore 1 μ l

7/3/01

PCR:

Template

pcr pm 3

0.5 μ l

1 μ l

control

5 μ l

5 μ l

5 μ l

dNTP

primer

1 μ l

1 μ l

1 μ l

buffer

5 μ l

5 μ l

5 μ l

Tag

0.5 μ l

0.5 μ l

0.5 μ l

H₂O

38 μ l

37.5 μ l

38.5 μ l

primer dilut.

95 μ l

H₂O

25 μ l

T7P1

25 μ l

CL dam M20

568 8362-118	Jun 12 2001
GENOSYS	568 8362-118
T7P1	
5'-GGATCCTGGGTTATCCCC	
TGATT	
1361.6 μ g	Tm=69.7°C
40.400	33.7 μ g/OD
195.0 nmol	MW=6981

568 8362-119	Jun 12 2001
GENOSYS	568 8362-119
CLDamM20	
5'-TCTAGATGTTGCCAATGC	
CAGTGTG	
578.8 μ g	Tm=68.2°C
18.300	31.7 μ g/OD
75.5 nmol	MW=7672

GENOSYS	568 8362-118
568	

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Signed

Signed

8/15/01 Cont.

PCR4 XLIB electroporation...

Blotted plates with Hybond membrane
followed protocol for colony hybridization

Cross linked membrane - stored at -20°C ON

8/16/01

Colony Hybridization PCR4 XLIB

made probe using 3ul PCR-4 + Tnl H2O in ECL Kit

preheat hybridization buffer to 42°C

wet screen mesh + membranes

Roll + put in hybridization tube

pre hybridize with 5X SSC for 1 hour

use 30mls hybridization buffer

probe Boil 5"

Cool on ice 5"

10ul labelling reagent

10ul glutaraldehyde

10' 37°C

add 1ml hybridization buffer from tube - add buffer +
probe to tube 42°C ON

ridoux

8/17/01

Colony Hybridization Cont.

preheat 1° wash buffer to 42°C

discard Hybridization buffer ~~Add 5X SSC~~

Add 50ml 5X SSC return to Oven 5min

discard 5X SSC add 1/3 vol of tube w/ 1° Wash Buf

Return to Oven for 20min

Wash again in 1° wash buff for 10min 2X

remove blots from tube place in glass container

Cover w/ 2X SSC

Shake @ RT for 5min X 2

Detection

8ml/membrane

incubate 1min

place on Saran

13mls #1

13mls #2

3membranes

Read and Understood By

Signed

Signed

Notebook Number: _____

Date: _____

9/19/01 Cont.

Culture reached 260 KU after 4 hrs incubation at 37°C

- (2) Put cultures into SS34 tube
- (3) Incubate on ice 15"
- (4) CF 4°C 5000 xg (6500rpm) 15"
- (5) Wash 2X w/1mM HEPES buffer (pH 7.0)
- (6) resuspend pellet in 10% glycerol to $1/12$ their
Original vol = 2mls
- (7) CF 5000xg 10"
- (8) resuspend in remaining glycerol solution after
supernatant is decanted
- (9) Flash freeze in $\text{EtOH} + \text{dry ice bath}$
- (10) Freeze at -80°C

P.mult 11039 is highly encapsulated - forms a very soft pellet
I lost the pellet in the 1st wash step, so more cells will
have to be prepared

Electroporate PLS88 into P.mult 1069

1ul PLS88 plasmid prep (in plasmid box spot G9)

40ul 1069 comp cells

2.5 KV
200 Ω
25 μF
 $T_c = 5.12 \text{ msec}$

Incubate 37°C 1 hr in LB

plate 10^0 , 10^{-1} , 10^{+1} 100ul on LB strep 37°C

10^{+1} = after plating 10^0 + diluting 10^{-1} , cf culture + resuspend pellet in
100ul media - plate 100ul

RESULTS
 10^{+1} 134 colonies
 10^0 2
 10^{-1} 0

9/25/01 Tuesday

Pass colony from R6E12 #4 blot - do a plasmid prep
Electroporate pLS88 into P.mult 11039

1 ul pLS88 plasmid prep (in plasmid box spot 59)

40 ul 11039 comp cells (flash frozen in liq N₂)

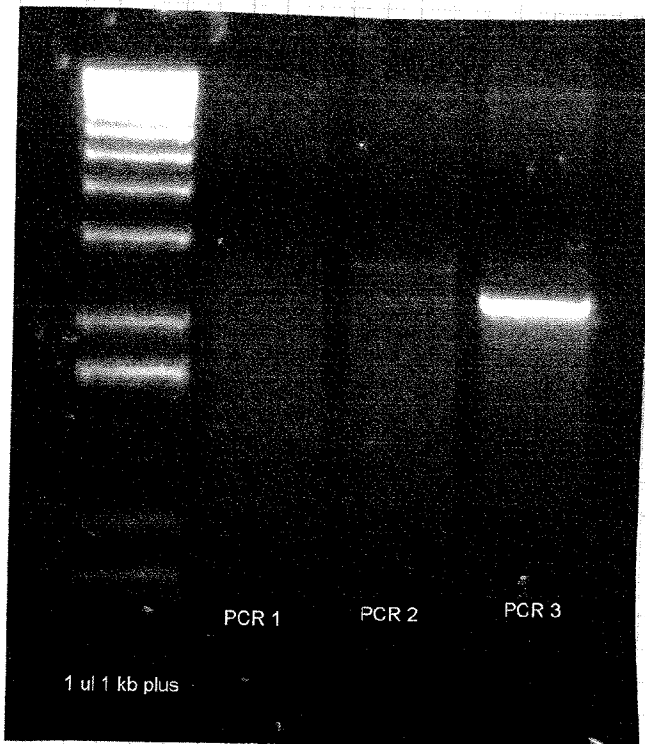
2.5KV
200 Ω
25 μ F
Tc = 5.16 msec

incubate 37C 1hr LB

Plate 10⁰, 10⁻¹, 10⁺¹ 100ul on LB strep 37°C

RESULTS

10⁺¹ 80
10⁰ 13
10⁻¹ 1



See pg 35 for PCR rx setup

Annealing temp = 48°C

Extension time = 30 sec

PCR rx did not work
again

0.7% agarose 3.5ul gel star
5ul of PCR Rx loaded

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Signed

Signed

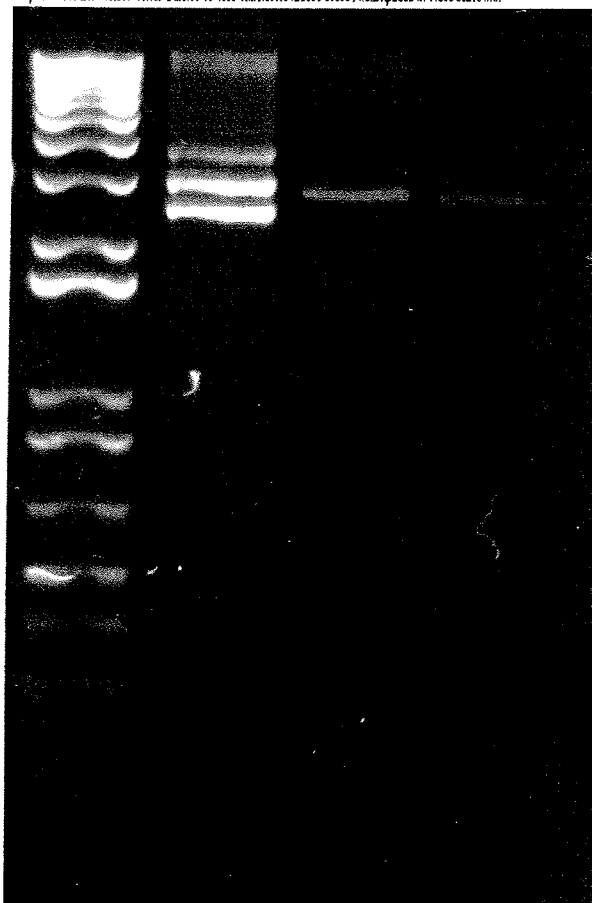
Notebook Number: _____

Date: _____

9/26/01 Wednesday - Gone to Memphis
 ML started 5ml culture of ~~RoEi2~~ ^{RoEi2} Ed 9 +
 9/27/01 Thursday

Did a Qia spin mini prep of RoEi2
 Eluted in 30ul EB

Exp: 1/30 Sec B:0 W:255 6:10:55 Date:00-00-1993 Time:00:00 ID:0000-00000 FileID:PLS88 in 11039 082701.tif



1ul
1kb
plus

RoEi2
1ul

PLS88
in
P.mult
11039

RoEi2 looks good
 Pass to a new plate
 freeze next week

PLS88 in 11039 looks
 good -

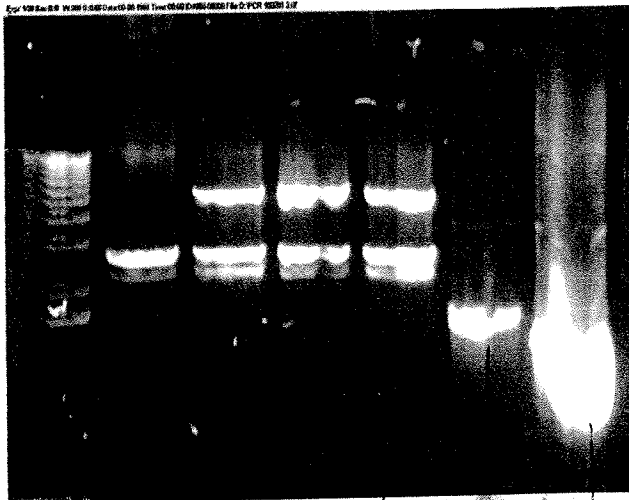
it work- we were
 able to electroporate
 into P.multocida
 Comp cells



PCR pCLpm3 T7P1 + CLdamM20 092801

Sample #	1	2	3
pCLpm3 8/29/01 BYR	0.5	1	0
dNTPs	5	5	5
T7P1 + CLdamM20	1	1	1
Taq	0.5	0.5	0.1
Taq Buffer	5	5	5
Water	38	37.5	38.
TOTAL Rx amt.	50	50	
Annealing temp	67.5 C		

10/2/01



See pg 41 pcr
Set up

PCLpm3
1ul

10ul cut out of gel = 0.4177 gm
put in Ref over night

10/3/01

Elute DNA from gel with Qiagen gel extraction Kit

cut gel fragment in 1/2 and put in 2 tubes

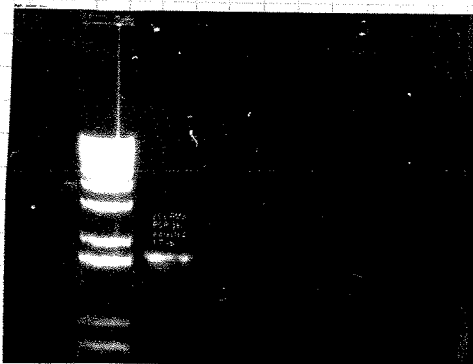
Added 600ul / tube of QG

Added 200ul isopropanol

eluted each column w/ 30ul EB into same tube = 60ul
total

Called pCLpm3 PCR

Loaded 5ul of pCLpm3 PCR on 0.7% agarose gel 3.5ul gel stain
100 Volts



Read and Understood By

Signed

Signed

Date: _____

10/3/01

Electroporated 40 μ l *P. multocida* 11039 Comp cells w/no DNA

Recover 1 hr in LB broth 37 C

plate 100 μ l 10^0 , 10^{-1} , 10^{-2} on LB
grow at 37 C O/NShould have plated on LB strep -
threw plates away

2.5 KV

25 μ l200 Ω $T_m = 5.16 \text{ msec}$

Before Digesting pCLpm3 PCR 1.7 Kb frag we need

more PCR product b/c XbaI has to be deactivated w/

EDTA (heat does not work) + then the DNA is ppt

out. DNA is lost in ppt so I need to amplify
my PCR product

PCR pCLpm3 1.7 kb T7P1 + CLdamM20 10/3/01			
Sample #	1	2	3
pCLpm3 PCR 1.7 kb	0.5	1	0
dNTPs 1 mM each	5	5	5
T7P1 + CLdamM20	1	1	1
Taq (7/14/01)	0.5	0.5	0.5
Taq Buffer	5	5	5
Water	38	37.5	38.5
TOTAL Rx amt.	50	50	50
Annealing temp	67.5 C		
extension time	30 sec		

run 0.5 μ l PCR rxns on gel

PLS88 digest BamHI/XbaI

10ul PLS88 plasmid prep 9/19/01
 1ul XbaI
 1ul BamHI
 2ul Multicore Buffer 10X
 6ul H₂O

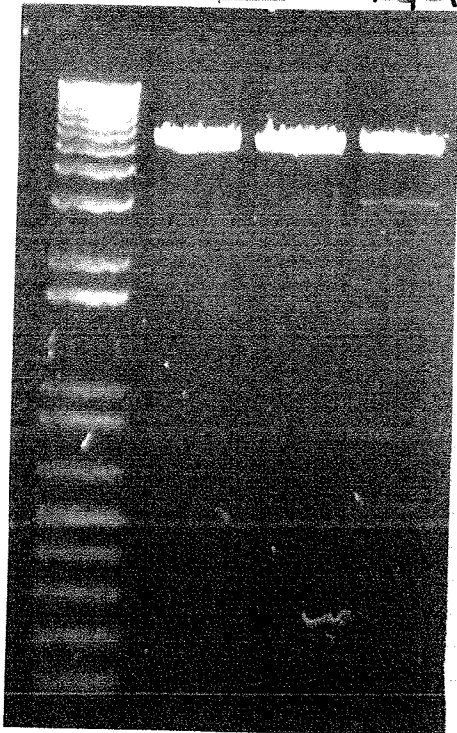
20ul

Digest 37C 4hrs Start 10:00am

Run 0.5ul of pCLpm3 1.7Kb PCR Rx run 10/3/01 on
 0.7% agarose gel 3.5ul gel star

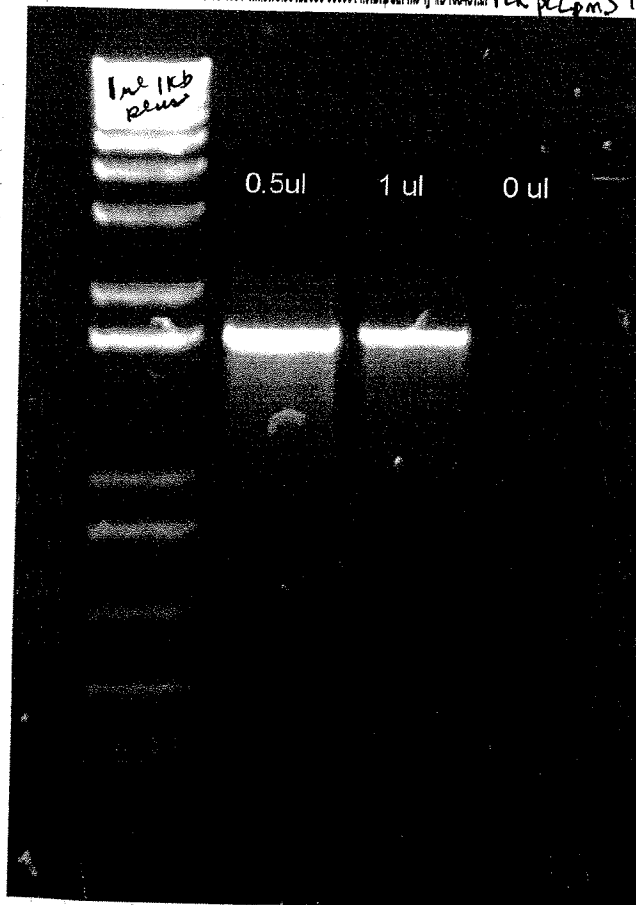
PLS88 BamHI XbaI digests

done 9/24/01



1ul 1ul 1ul 1ul
 1kb Bam Xba
 plus HE I
 plus

Exp: 130 Ser: B.0 W: 265 G: 0.65 Date: 00-00-1993 Time: 00:00 ID: 0000-00000 File: D:\bcl\PAR 17 kb 100401.tif PCR pCLpm3 1.7Kb



Read and Understood By

Signed

Signed

Date: _____

10/4/01

Digest pCLpm3 1.7Kb frag PCR Rx 1 (from 10/3/01)

10ul	DNA	
1ul	Xba I	
1ul	Bam HI	37 C
2ul	MultiCore 10X Buffer	4 hrs
6ul	H ₂ O	
<hr/> 20ul		

To pLS88 + pCLPM3 1.7Kb Bam HI Xba I digests

Add	0.5ul	0.5M EDTA	to deactivate Xba I
Add	2ul	3M Na Acetate	(1/10 vol)
	44ul	ETOH	44 (2 vols)

freeze -20 over night to ppt DNA.

Start 2 1L BHI Ap cultures of R6E12 for LPS prep

Add 2.5mls of ON culture per Liter

Add 500ul 200mg/ml Ap = 100ug/ml final vol
1/2 normal conc. used

grow ON 26°C Shaking

10/5/01

Cf. R6E12 cultures in 250ml bottles on GST rotor

6000rpm for 15min at 4°C

freeze pellets at -20 C.

Do LPS prep next WK

10/5/01

cf. ppet DNA pLS88 + pCLPM3 1.7kb
 resuspend in 10 μ l TE. put at 4C to go into
 Suspension

10/8/01

run 1ul pLS88 + pCLPM3 1.7kb on gel
 ✓ on gene spec first

Sample	Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
pCL PM 3 1.7 kb	dsDNA	0.655	1.536	0.818	0	1.87775	0.0768 ug/ μ L	1	50	104.32
	dsDNA	0.707	1.592	0.85	0	1.87294	0.0796 ug/ μ L	1	50	104.05
pLS88 Bam HI Xba I	dsDNA	1.565	3.982	2.369	0	1.68088	0.1991 ug/ μ L	1	50	93.38
	dsDNA	1.987	4.493	2.85	0	1.57649	0.22465 ug/ μ L	1	50	87.58

pCLPM3 76.8 ng/ μ lpLS88 211.9 ng/ μ lLigation

pLS88 4.5 kb

pCLPM3 1.5 kb

pLS88 1ul 200ng
 pCLPM3 7.5ul 570ng
 Ligase buff 1ul
 Ligase 0.5ul
 10ul

14°C overnight

10/9/01 Tuesday

Heat inactivate ligation 65°C 15"

Read and Understood By

Signed

Signed

PCR 93146 Ei OPS primers 10/17/01

Sample #	1	2	3	4
93146 genomic prep 7/28/99	2	2	2	2
dNTPs 2.5mM	10	10	10	10
Ei OPS PCR U 2 + Ei OPS PCR R 2	1.2	1.2	1.2	1.2
EXL polymerase	1	1	1	1
DMSO	0	0.5	1	1.5
Stabilizing Soln	1	1	1	1
10 X Buffer	5	5	5	5
Water	29.8	29.3	28.8	28.3
TOTAL Rx amt.	50	50	50	50
Use PCR EXL Pcr Program				
Annealing Temp	63			
PCR EXL primer				

Ran out of EXL polymerase
after 1st rx.

ran 1st rx
ordered more EXL pcr

Rxs 2-4 are in green ice box
- 20 chest freezer

gel picture pg 54

Make 2 5ml LBAP cultures of pFPV25 in DH10Bs

- 1 - plasmid prep
- 1 - freeze

Make 5ml culture of pLS88 isolates pos + Adj on LBstrep
positive colony → adjacent to
from ~~last~~ hybridization positive colony

Read and Understood By

Signed

Signed

Date: _____

10/18/01

Started cultures in wrong Antibiotics
 started pFPVas in LB strep - pLS88 in LBAP Opps -

Start new 5 ml cultures
 pFPVas in LB AP
 pLS88 in LB strep

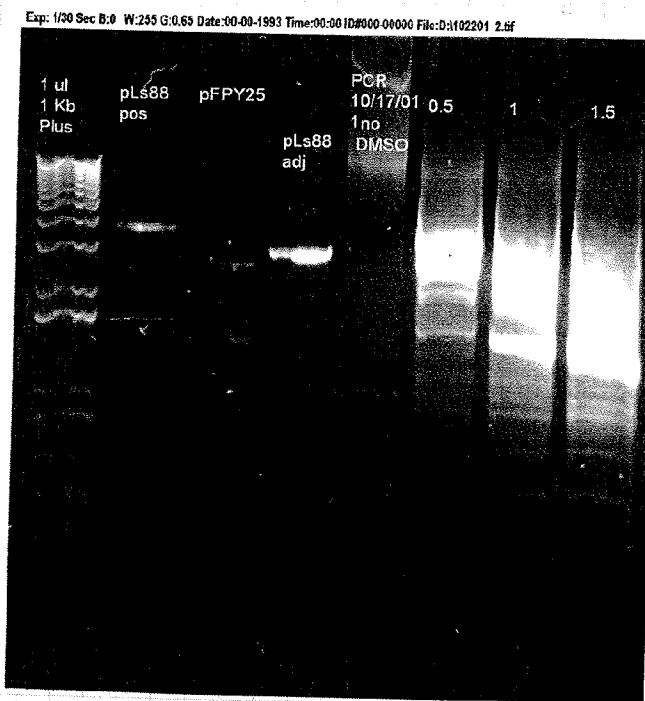
10/19/01

Froze 1 pFPVas culture (Ask BYX where it is)

Spin down other pFPVas culture + both pLS88 cultures
 (pos + adj) - freeze pellets for plasmid preps

10/22/01

made Qiaspin plasmid preps of
 pFPVas, pLS88 positive, pLS88 adjacent



Loaded 1ul of
 pLS88 pos
 pLS88 adj
 pFPVas

Loaded 5ul of
 PCR product

pass both pLS88 colonies
 to a new LB strep plate

Read and Understood By

10/23/01

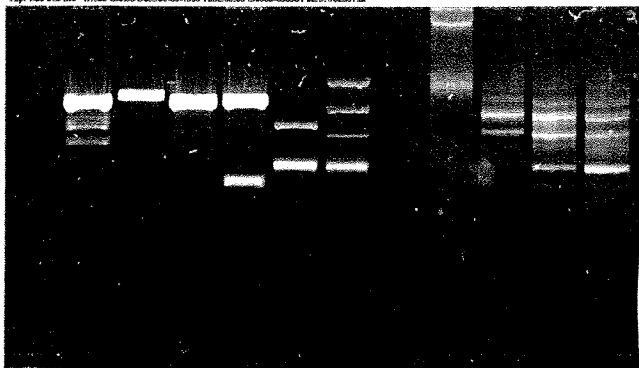
Digest PLS88 pos + Adj plasmid prep (10/22/01)

Cutcut
 insert
 PLS88 pos 4 ul
 Bam HI 1 ul
 Xba I 1 ul
 multicores 2 ul
 H₂O 12 ul
 PLS88 Adj 4 ul
 BamHI/XbaI cuts
 the insert out
 20 ul
 load 10 ul on gel

linearize
 PLS88 pos 2 ul
 EcoRV 1 ul
 Buff D 1 ul
 H₂O 6 ul
 PLS88 Adj 1 ul
 EcoRV 1 ul
 Buff D 1 ul
 H₂O 7 ul
 10 ul
 load 10 ul on gel

linearize
 DNA
 PFPV25 5 ul
 EcoRV 1 ul
 Buff D 1 ul
 H₂O 3 ul
 PFPV25-
 pst 5 ul
 Buff H 1 ul
 H₂O 1 ul
 10 ul
 load 10 ul on gel

Fig. 100 Sec B0 W:255 C:0.55 Date:01.09.1993 Time:07:20 E14000.00000 File:01102201.tif



1 kb plus
 did not show
 up

- 1) PLS88 adj BamHI/XbaI
- 2) PLS88 pos EcoRV
- 3) PLS88 Adj EcoRV
- 4) PLS88 pos BamHI/XbaI
- 5) pFPV25 pst I
- 6) pFPV25 EcoRV
- 7) PCR 10/17/01 No DMSO 2 ul DNA + 1 ul 6x dye
- 8) ↓ 0.5 ul ↓ + 3 ul H₂O
- 9) ↓ 1 ul ↓
- 10) ↓ 1.5 ul ↓

Read and Understood By

Signed

Signed

Date: _____

10/24/01

PCR pMBE1 Ei OPS primers 10/24/01				
Sample #	1	2		
From 9/17/00 in pMBE1 DNA Box	0.5	0.5		
dNTPs 2.5mM	10	10		
Ei OPS PCR U 2 + Ei OPS PCR R 2	1.2	1.2		
EXL polymerase	1	1		
DMSO	0	0.5		
Stabilizing Soln	1	1		
10 X Buffer	5	5		
Water	31.3	30.8		
TOTAL Rx amt.	50	50		
Use PCR EXL Pcr Program				
Annealing Temp	63			
PCR EXL primer				

Read and Understood By

PLS88 positive gene spec

Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
dsDNA	3.461	5.406	4.733	0	1.14219	0.2703 ug/uL	1	50	63.46
dsDNA	0.046	1.798	1.112	0	1.61691	0.0899 ug/uL	1	50	89.83
dsDNA	0.018	1.781	1.091	0	1.63245	0.08905 ug/uL	1	50	90.69
dsDNA	0.031	1.779	1.083	0	1.64266	0.08895 ug/uL	1	50	91.26
dsDNA	0.3	2.076	1.421	0	1.46094	0.1038 ug/uL	1	50	81.16
dsDNA	0.732	2.531	1.855	0	1.36442	0.12655 ug/uL	1	50	75.80
dsDNA	0.538	2.393	1.707	0	1.40187	0.11965 ug/uL	1	50	77.88
dsDNA	0.37	2.2	1.522	0	1.44547	0.11 ug/uL	1	50	80.30

pls88 plasmid prep made 10/24/01 $\text{avg} = 124.8 \text{ ng/uL}$

Seq pls88 w/ CLPM1-PL 3.2 pmol \rightarrow p.29 bKV + w/T3 3.2 pmol \rightarrow pg 27 bKV

PLS88 4ul

TRR 8ul

primer 2ul

H₂O 6ul

20ul

#1 CLPM1-PL

#2 T3

Sequence should contain DAM gene sequence

CLPM1-PL primer sequence data is trashy - but

I blasted a segment of it and it is the DAM gene sequence

Redigest pFPV25 5ul

Buff H 1ul

H₂O 3ul

ECORI 1ul

~~ECORI~~ 1ul

Pst I 1ul

10ul digest

pFP125 5ul

ECORI 1ul

Pst I 1ul

Buff H 2ul

H₂O 10ul

20ul

Read and Understood By

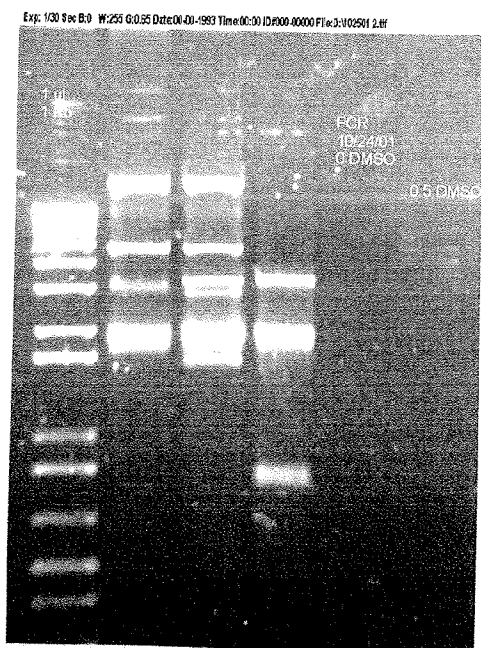
Signed

Signed

Notebook Number: _____

Date: _____

10/25/01



0.7% agarose 3.5ul gel star

- 1) 1ul 1 Kb plus
+ 4ul H₂O
+ 1ul 6x dye
- 2) pFPV25 EcoRI 10ul + 2ul
- 3) pFPV25 pstI 10ul
- 4) pFPV25 EcoRI/pstI 10ul
- 5) PCR pmBE1 0.5ul No DM
- 6 " " " 0.5 DMSO
+ 1ul 6x dye
3.5ul H₂O

Freeze pLS88 pos ~~XXXXXXXXXX~~
Start 5ml LB strep

Electroporate pLS88 pos into P. mult 11039 comp cells

2.5 kV
200 Ω
2.5 μ F
TC = 5.08ms

40ul P. mult (All D4 9/21/01)
1ul pLS88

Recover 1 hr LB strep

plate 10^{+1} on LB strep
incubate 37°C ON

10/26/01

Exhibit A Page 46

Make Qiaspin plasmid prep of pFPV25 in DH10βs
Elute 30ul EB

Digest w/ EcoRI / Pst I (same as yesterday)
digest 5 hrs

run on gel w/ undigested plasmid

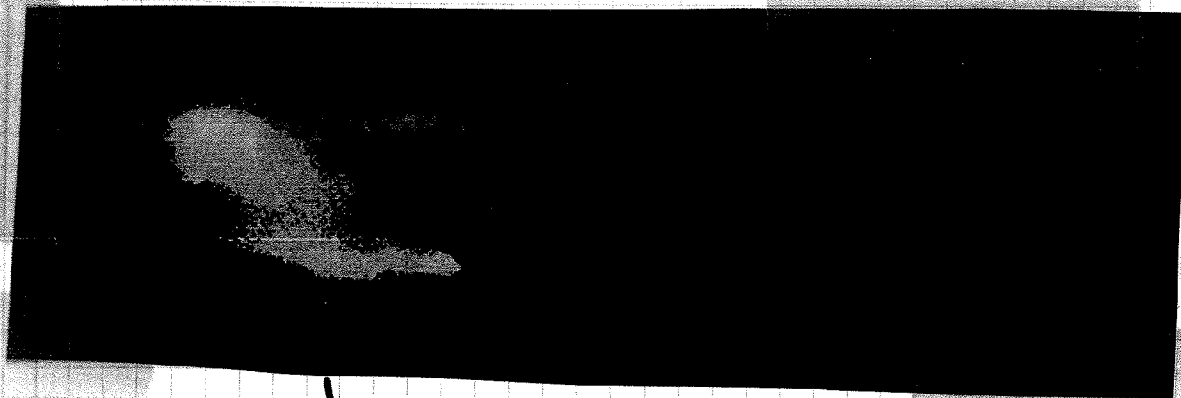
Freeze Pls 88 pos now called pls 88 DAM
it is in XL1B MRF'

ALL D5-9 LBstep + 20% glycerol

Put plate from electroporation in ref over weekend -
Monday pick 4 big colonies + plate on LB step

Started BHI Ap plate cultures of 93146 WT Lux +
93146 R⁻ Lux

Imaged fish immersion → ip injected w/ 93146 WT Lux
on Night Owl.



→ Luminescent area

Read and Understood By

Signed

Signed

PLS88 positive gene spec

Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
dsDNA	3.461	5.406	4.733	0	1.14219	0.2703 ug/uL	1	50	63.46
dsDNA	0.046	1.798	1.112	0	1.61691	0.0899 ug/uL	1	50	89.83
dsDNA	0.018	1.781	1.091	0	1.63245	0.08905 ug/uL	1	50	90.69
dsDNA	0.031	1.779	1.083	0	1.64266	0.08895 ug/uL	1	50	91.26
dsDNA	0.3	2.076	1.421	0	1.46094	0.1038 ug/uL	1	50	81.16
dsDNA	0.732	2.531	1.855	0	1.36442	0.12655 ug/uL	1	50	75.80
dsDNA	0.538	2.393	1.707	0	1.40187	0.11965 ug/uL	1	50	77.88
dsDNA	0.37	2.2	1.522	0	1.44547	0.11 ug/uL	1	50	80.30

pls88 plasmid prep made 10/24/01 $\text{avg} = 124.8 \text{ ng/uL}$

Seq pls88 w/ CLPM1-PL 3.2 pmol \rightarrow p.29 bKV + w/T3 3.2 pmol \rightarrow pg 27 bKV

PLS88 4ul

TRR 8ul

primer 2ul

H₂O 6ul

20ul

#1 CLPM1-PL

#2 T3

Sequence should contain DAM gene sequence

CLPM1-PL primer sequence data is trashy - but

I blasted a segment of it and it is the DAM gene

sequence

Redigest pFPV25 5ul

Buff H 1ul

H₂O 3ul

ECORI 1ul

~~ECORI~~ 1ul

Pst I 1ul

10ul digest

pFP125 5ul

ECORI 1ul

pst I 1ul

Buff H 2ul

H₂O 10ul

20ul

Read and Understood By

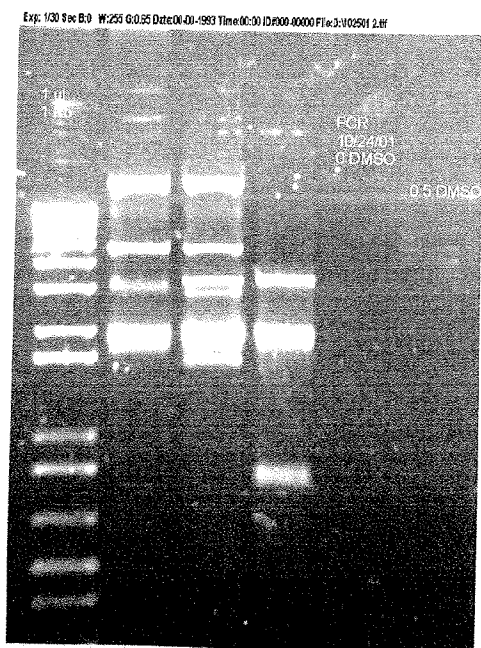
Signed

Signed

Notebook Number: _____

Date: _____

10/25/01



0.7% agarose 3.5ul gel star

- 1) 1ul 1 Kb plus
+ 4ul H₂O
+ 1ul 6x dye
- 2) pFPV25 EcoRI 10ul + 2ul
- 3) pFPV25 pstI 10ul
- 4) pFPV25 EcoRI/pstI 10ul
- 5) PCR pMBE1 0.5ul No DM
- 6 " " " 0.5 DMSO
+ 1ul 6x dye
3.5ul H₂O

Freeze pLS88 pos ~~XXXXXXXXXX~~
Start 5ml LB strep

Electroporate pLS88 pos into P. mult 11039 comp cells

2.5 kV
200 Ω
2.5 μ F
TC = 5.08ms

40ul P. mult (All D4 9/21/01)
1ul pLS88

Recover 1 hr LB strep

plate 10^{+1} on LB strep
Incubate 37°C ON

10/26/01

Make Qiaspin plasmid prep of pFPV25 in DH10 β s
 Elute 30ul EB

Digest w/ EcoRI / Pst I (same as yesterday)
 digest 5 hrs

run on gel w/ undigested plasmid

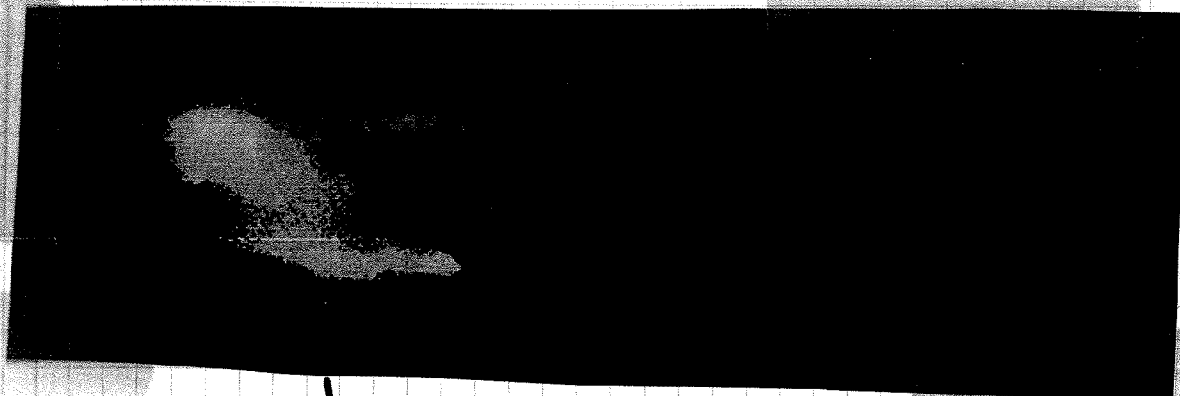
Freeze plasmids now called plasmids
 it is in XL1B MRF'

ALL D5-9 LBstep + 20% glycerol

Put plate from electroporation in ref over weekend -
 Monday pick 4 big colonies + plate on LB step

Started BHI plate cultures of 93146 WT Lux +
 93146 R⁻ Lux

Imaged fish immersion \rightarrow ip injected w/ 93146 WT Lux
 on Night Owl.



\rightarrow Luminescent area

Read and Understood By

Signed

Signed

electroporation. grow on LB strep

Notebook Number: _____

Date: _____

11/5/01

pour 12% / 4% SDS-PAGE gel
gel stayed in buffer in 4°C ON

Start 5ml LB strep cultures of P. mult 11039 pLS88 From 10/31
11/6/01

Samples to run on gel

LPS from R6Ei2 dilute - 1/1 in sample Buffer

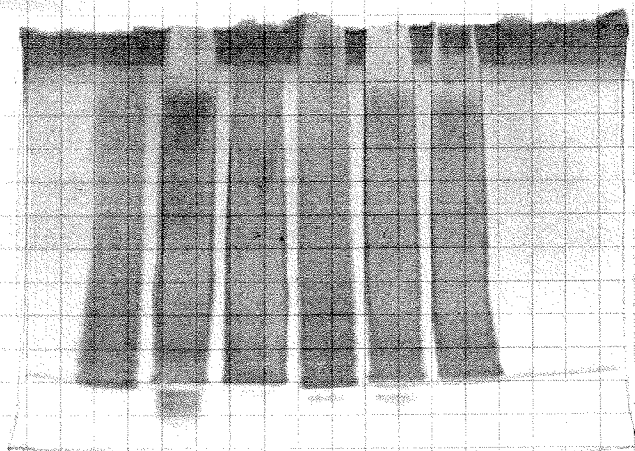
and take 10ul (1mg/ml) + 15ul H₂O + 25ul Buffer = 10ul

LPS Salmonella typh 1/1 of 2mg/ml in H₂O 1/1 in Buffer

Lane 1	Sal typ	}	Not Denatured
2	R6Ei2 1/1		
3	R6Ei2 10ul		
4	Sal typ	}	Denatured
5	R6Ei2 1/1		
6	R6Ei2 10ul		

loaded 10ul / well Ran at 100volts

Silver stain for LPS



Made Qia spin plasmid preps of P. mult 11039 pL588
4 colonies

11/7/01 Run 12%/4% SDS PAGE gel
Same as 11/6/01

- 1) Sal. typ 1/1 of 2mg/ml in H₂O 1/1 in Buffer - Load 10ul
- 2) R6E12 1/1 in buffer
- 3) R6E12 10ug
- 4) wt 93146 LPS 10ug

Electrophoresed at 100V

transferred to Nitrocellulose 100 volts 1 hr

Blocked membrane overnight 5% NFM in PBS

Our Ed 9 was contaminated - so was Bobbica's

DH103a pFPR25 frozen -80C All F1-5

tube #1 used to start a 5ml LBap culture

+ put on LBap plate

P. mult 11039 pL588 #1 Start 5ml LB strep culture to
freeze → see off pic next page

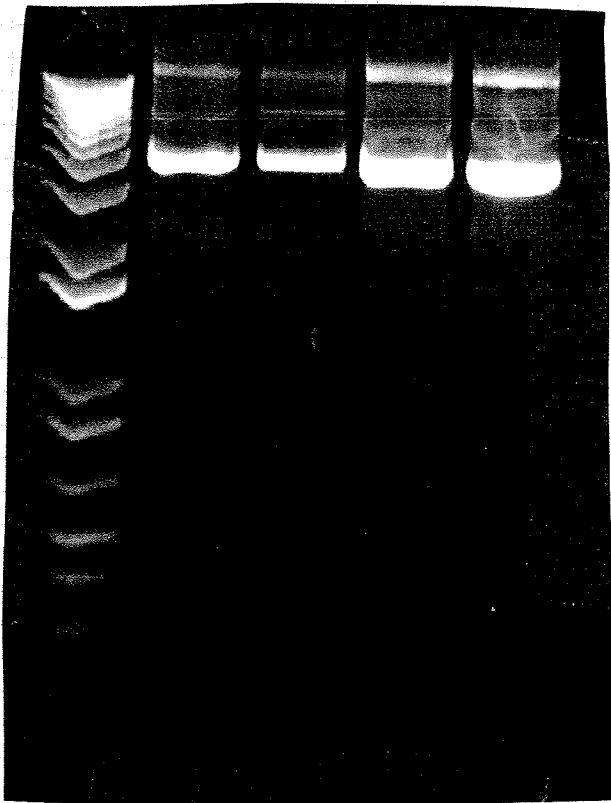
Read and Understood By

Signed

Signed

Notebook Number: _____

Date: _____



p. mult 11039
 Dan gene. pLs 88
 plasmid preps
 from 11/6/01

1ml
 1Kb plus

11/8/01

Western Blot R6E12

There is No Ed 9 to use as 1^o Ab so

1^o used α Ed 1ct 4383 polyclonal CF serum 1/1600
 + α Ed 1ct 93146 wt polyclonal CF serum 100ul added
 to 1/1600 dilution

Incubate RT 1 hr shaking
 Wash 3X PBS TW20

2^o 9E1 1/4 1 hr RT shaking
 Wash 3X PBS TW

30 goat & mouse Ig AP 1/1500 1 hr RT Shaking
 Develop BCIP/NBT 20 mes

it did not work
 will have to repeat
 when Bobbie has more
 Ed 9

1 2 3 4

Frog pasturella multocida 11039 p1588 (DAM gene)
 LB + 20% glycerol
 -80 C Box A11 G1-5

Digest p1588 p.mult 11039 #1

BamHI/XbaI

DNA 4ul
 BamHI 1ul
 XbaI 1ul
 multicore 2ul
 H₂O 12ul
20ul

BAM HI

DNA 4ul
 BamHI 1ul
 multicore 1ul
 H₂O 4ul

Digest 37°C
 for 2 hrs approx
 run on gel tomorrow

Plasmid prep pFV25 in DH10βs
 Eluted in 30ul set next pg for gel pre

Read and Understood By

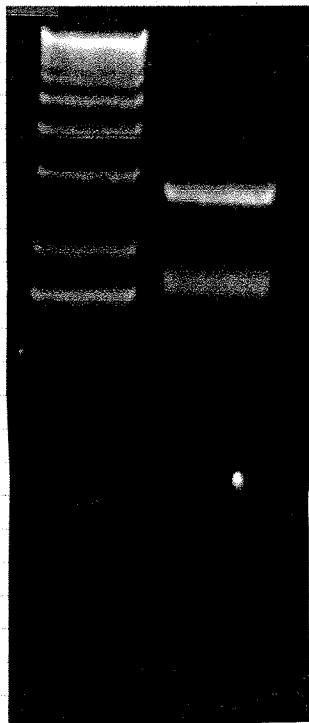
Signed

Signed

Notebook Number: _____

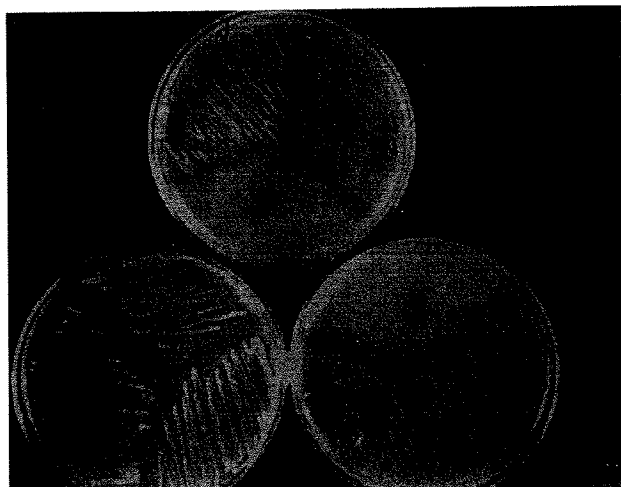
Date: _____

11/8/01



pFPR25

— do not know what this band is

next run on gel and extract top
band; electroporate into XL1B

Exp: n/a Bin: 1x1 Gain: 1.0 B.O.W: 6535 G.O.55 N.O Date: 11/1/2001 Time: 11:14:39 am ID#0 File: Untitled

Neg Controls
14 + 15

22 = 93146 Lux

4d post
Red fluorescence
immersion93146 Lux
recovered from
fish immersion
infectedBacteria was
cultured
4 days
post
immersionLux plasmid
is still
stable
in vivo

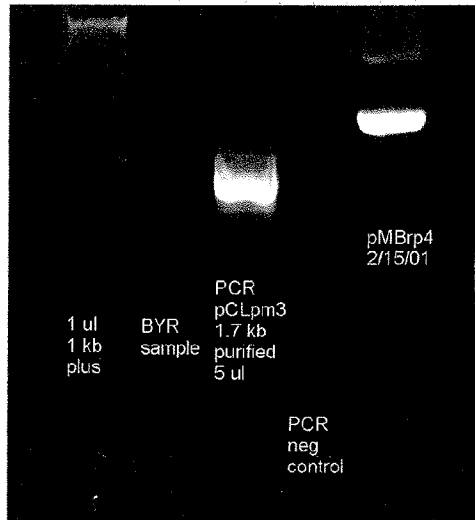
PCR pCLpm3 1.7 kb T7P1 + CLdamM20 purified
01/16/02

Sample #	1	2	
pCLpm3 PCR 1.7 kb purified	0.5	0	
dNTPs 1mM each	5	5	
T7P1 + CLdamM20	1	1	
Taq (7/16/01)	0.5	0.5	
Taq Buffer	5	5	
Water	38	38.5	
TOTAL Rx amt.	50	50	
Annealing temp	67.5 C	program m1 under mark	
extension time	30 sec		

1/18/02

run 5ul on gel of PCR rx

5ul DNA
1ul 6x Dye



measured pMBRp4 on gene spec - got a low reading of 11.8 ng/ul
so ran 1ul on gel -
conc on gel looks like = 100 ng/ul

digest w/ Ban HI Hind III

double digest

pMBRp4	4 ul	4
enzyme	0.5 ul	0.5 each
10X buff E	1 ul	1
H2O	4.5 ul	4

digest from 2:00 - 5:00 37°C

Read and Understood By

Signed

Signed

2/13/02

Ligation: To End Conversion Rx mix add:

- (1) 1ul pL588 vector (pL588 EcoRV digest)
1ul ligase
 - (2) + control
1ul pT7 Blue vector
1ul ligase
 - (3) - Control
1ul pT7 Blue vector
1ul ligase
- 22°C 15"

~~plate~~

Transformation:

Add 1ul ligation Rx to 1 tube of Nova Blue Comp cells

Incubate on ice 5min

heat shock 30 sec in 42°C water bath

Incubate on ice 2min

Add 250ul RT SOC media

plate

(+) + (-) Controls 50ul @ 1/10 (5ul in 45ul H₂O)
on LB S-gal

exp 50ul @ 1/10 on LB strep

2/14/02

Colonies from ligation were very small first thing
this AM.

Read and Understood By

Signed

Signed

Notebook Number: _____

Date: _____

3/14/02 Thursday

PCR pCLpm3 1.7 kb T7Sal I P2 + CLdamECoRV
3/14/02

Sample #	1		
pCLpm3 PCR 1.7 kb purified (1/23/02)	0.5		
dNTPs 1mM each	5		
T7Sal IP2 + CLDamECoRV	1		
Taq (7/16/01)	0.5		
Taq Buffer	5		
Water	38		
TOTAL Rx amt.	50		
Annealing temp	55		
extension time	30 sec		

used wrong template
 repeat using
 pCLpm3 plasmid prep
 8/29/01 red box G8

15 7389-065 Mar 7 2002
 GENOSYS
 CLDamECoRV
 5'-CTGTTTGGATGATGCTGCT
 TTICA
 3418g 10.000 47.3nmol
 Tm=60.3°C 34.2ug/DD MW=7225

15 7386-065 Mar 7 2002
 GENOSYS
 T7SalIP2
 5'-GGGCTGATTGCTGGGACA
 TTTCA
 5143ug 15.400 95.1nmol
 Tm=62.4°C 33.4ug/DD MW=5410

reconstituted primers

2.5ul each + 95ul H₂O

made new 1mM dNTPs

40ul 10mM mix to 60ul H₂O

plated pLSS8 from frozen stock A3B4 put on LB strep at
 37C MLL will take out of incubator + put at 4 C
 tomorrow

Notebook Number: _____

Date: _____

4/16/02

Hybridize colonies from Miss RCPm3 ligaton
follow protocol pg. 50

probe

PCR prod green PCR box A9 - PCR rx 3/21/02
pg. 94 bkt = 5.35 ng/ul

Cleanup PCR prod using microcon columns -
follow directs in kit

removes primers & dNTPs

Read on Gene Spec -

Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor
dsDNA	0.709	0.366	0.224	0	1.63393	0.0183 ug/uL	1	50
dsDNA	0.736	0.655	0.497	0	1.31791	0.03275 ug/uL	1	50
dsDNA	0.492	0.945	0.804	0	1.17537	0.04725 ug/uL	1	50
dsDNA	0.551	0.997	0.861	0	1.15796	0.04985 ug/uL	1	50

There are crazy readings - gene spec is
not good for measuring DNA of PCR prod.

MLL said use 3ul in probe

Remained of cleaned-up PCR prod is in Pink Box H8

Hybridize ON 42°C

4/17/02

Developed Hybridization

1" film almost had no image
10" very light image
1 fr OK

Plate 1

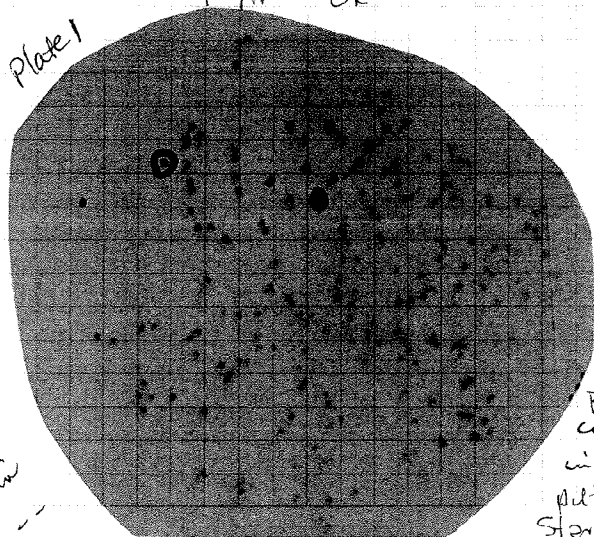
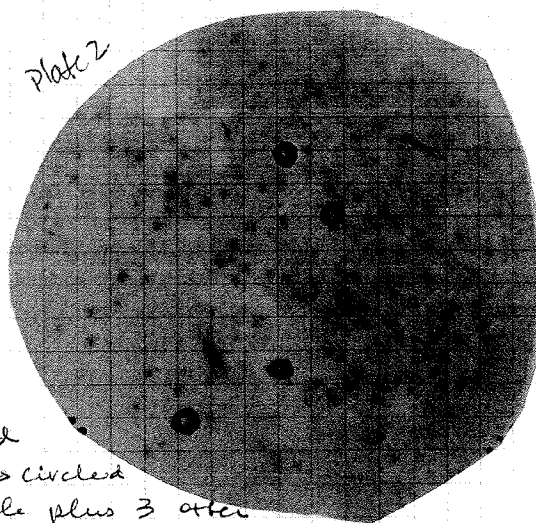


Plate 2



passed
colonies circled
in purple plus 3 other
put on LB strep
Start Sml Ubaton & Bactant & Nitrofurantoin & BV

plates in
1 -

5/13/02

cycle seq PGEM 32 test plasmid w/ M13 primer from A61
 run on seq. to create matrix file

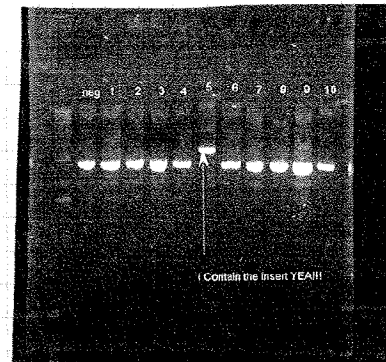
5/14/02 Tues

Clones from Electroporation grew well
 grow 2ml LB strep cultures
 pick colonies w/ sterile toothpick
 use 1ml tomorrow to do plasmid preps to ✓ for insert
 grow 1 culture from Neg Control

plsgg pcpm3	100ul	= 150	53
	50ul	26	5

Neg Con (plsgg)	100ul	21	14 35
No insert	50ul	14	11

pick 10 ligation colonies
 1 neg con colony



5/15/02 Wed

all 2ml cultures grew
 used 1ml to do Qimprep spin plasmid preps
 eluted in 30ul EB

Run 3ul DNA on gel
 1 ul 6x Dye
 2 ul H₂O
 6ul

It Finally
 worked !!

#5 & Neg
 Plasmid preps
 in yellow box
 A5 + A6

For Vaccine trial:

plated 93146 R₆ from 6/19/01 A4G5 -80C on Blood
 grow at 26°C 48hrs

Reconstituted Intervet Edict vaccine - filled bottle 2/3 full w/ sterile H₂O

plate 1 loop full on Blood grow at 26°C 48hr
 put 50ul into 5ml BHI to freeze back grow at 26°C shaking over night

1 750ml flask BHI for Intervet Vaccine
 4 750ml flask BHI for R₆

Read and Understood By

Signed *Michelle Se*

Signed

For Vaccine trial cont:

Took 5mls of heat killed Ed 1ct (BVR's prep 4/12/02)

Tared tube empty
add 5mls

at 12000rpm 10"

Aspirate Sup

Re weigh tube = 0.0499 gm

Lyophilize overnight + re weigh

resuspended pellet in 2mls H₂O

Transfer to a pre-weighed
15 ml tube = 6.6018 gm

Freeze

Lyophilize

15 ml tube Dry wt = 6.6064

Dry wt of pellet = .0046

Digest pLs 88 pCLpm3 + clone (#5) + Neg plasmid preps

w/ Sal I + Cla I single digests

2ul DNA

1ul Buffer

0.5ul Enzyme

6.5 ul H₂O

10ul digest

ABI MaxA matrix stds came in. stds are 4 colors: Blue, Red, Green, yellow

mixed 1ul of each Standard with 12ul of deionized formamide
(Borrowed from Hanson's Lab). 1 tube per std = 4 tubes 1 each color

Heat to 95C for 2 "

Chill on ice

Sequence using Seq Run (250ul) E run module

injected each sample 3 times - this is done when the injection
list is filled out

Date: ~~11/5/02~~ 5/16/02

5/16/02 Thurs

Stats are still running on seq.

Weighed lyophilized pellet of heat ~~kill~~ prepare 2-750 ml cultures of heat killed Ed. 1ct - Monday

Run digests on gel. pLys8 pel p/n electroporate plasmid prep clone #5

Load entire Digest

6X dye is really conc. So only use 100ul

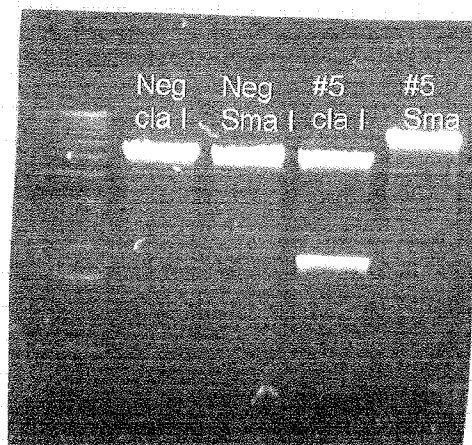
10ul DNA digest

1ul 6X Dye

1ul H₂O

12ul on gel

Lane 1 1ul 1kb
 2 Neg Cla I
 3 Neg Sma I
 4 #5 Cla I
 5 #5 Sma I



Electroporate: pLys dam 2 into 11039 p.mult. electrocompetent cells

1ul plasmid prep #5 (yellow box) A8

40ul 11039 Comp cells box A11 D2

Tc = 5.12

5.5 KV

200 Ω 25 μ F

Recovered 1 hr in LB 37C

plate 100ul neat on 2 LB Strep plates

Made 2 more 750ml broth cultures to grow 93146 wt #19

to heat ~~kill~~ Kill for vaccine trial

Read and Understood By

Michelle S.
 Signed

$37g \times 750ml = 27.75gms$ BH1

5/16/02 Thurs.

Froze

pls Dam 2 5vials 1ml/each in LB_{strep} + 20% glycerol
Aguavac-Esc 5vials 1ml/each in BHI to 20% glycerol
vaccine

5/17/02 Fri.

Staff Appreciation Day.

Nothing has grown yet on plates from Electroporation -

maybe need to be on BHI_{strep}
Maybe 11039 Comp cells Dead

Leave at 37 C until Tomorrow

Passed 93146 WT #19 to new blood plate

Start 5ml culture Sun

Took Rec + vaccine plates out of incubator
Leave at RT

Start 5ml BHI cultures Sat.

5/19/02 - Started 5ml BHI cultures

5/20/02 MM leave Mark Sick BYR started 2 750ml cultures
of ~~WT~~ WT #19

5/21/02 93146 WT Ed. let take 1ml from each 750ml culture
Combine in 1 tube

Streak for purity - plate is pure

plate 10^{-5} + 10^{-6} for colony counts

10^{-5} TNTC TNTC

10^{-6} 286 287 Avg 286.5

Heat Kill at 60°C for 3 hours
plate for Viability - NO growth at 48 hrs

7/24/02

Blotted plates on Nitrocellulose membrane

Block in 0.01M PBS + 5% NFM 2 hrs RT

1° Ab used Straight Ed 9 1 hr RT

3X wash PBS 0.01M + Tween

2° Ab goat anti mouse Ig(H+L) AP 1/1500 4°C Overnight

Set up Serum Killing Assay see pg 17 for plate setup

Serum HI at 56°C for 30 min

Read plate every 5 min for 1 hr

plate 15 μ l of 10^{-4} + 10^{-4} on BHI Ap

Colony Count Results (7/25/02)

E. coli 10^{-4}
27, 30, 47 10^{-7}
2, 2, 3assay worked
perfectly
 2.31×10^9
 1.56×10^9 7/26/02 WT 10^{-4}
35, 21, 26 10^{-7}
2, 5, 9 1.82×10^9
 3.56×10^9 RC 10^{-4}
44, 30, 35 10^{-7}
0, 3, 6 2.42×10^9
 2×10^9

Injected mice

100 μ l ip. of p.multocida WT 50, 100, 1000 2 mice per dose

" Dam 50, 100, 1000

" Dam 2 50, 100, 1000

Controls PBS 3 mice


 Signed

Read and Understood By

 Signed

24 CO₂ indicates mouse was euthanized to reduce suffering

Notebook Number: _____

Date: _____

7/25/02 Mortality from mice Exp.

9:00am WT 1000 2 mice 1st DOA 2nd CO₂ lethargic, Shivering, Flaccid, pilo-erection
 WT 50 1 CO₂ " " "
 Dam 1000 1 DOA
 Dam 100 1 DOA
 Dam 2 100 1 DOA

Notes:

2nd mouse in WT 50 Ht is very lethargic, doesn't respond to being handled
 2nd mouse in Dam 1000 is lethargic + shivering (due to fever?)

11:30am WT 100 CO₂ lethargic, Fever (shivering) does not respond to handling
 Dam 1000 1 DOA
 Dam 50 1 DOA

Notes:

2nd mouse in Dam 50 is sick - Shivering, non-responsive, piloerection

1:30 No more

4:45 WT 50 1 DOA
 Dam 50 1 DOA
 WT 100 1 DOA

All mice from Trt Groups WT 1000, WT 100, WT 50, Dam 1000 + Dam 50 are Dead

Dead mice were opened ventrally, the chest was spread open
 : they were stored in Sample cups w/ 10% Buffered formalin

7/26/02

9:00am Dam 100 1 DOA

All mice from trt Group Dam 100 are Dead

7/27 + 7/28 Dan Scruggs ✓ mice

Mon 7/29/02

Start WT, R6 + E. coli Lux cultures on BHI + 5ml BHI broth

Mice - Am all Trt groups are active + Alert
 pm " " " " "

8/28/02

Colony counts- Serum Killing Assay *E. coli* Lux10⁻⁷5
5
5 1.6×10^{10} CFU/ml10⁻⁶35
51
25 3.7×10^{10} CFU/ml

8/29/02

Re lux + WT Lux Colony Count plates were
Contaminated too badly to count

9/5/02

Inject mice

BYR prepared *p. multaxida* 11039 WT + Dan2 for
injections

1 extra mouse un.injected

5 mice / dose 9 treatments

20 - 4 doses WT 5, 10, 50, 100

20 - 4 doses Dan 2 10, 50, 100, 500

5 Control mice - PBS

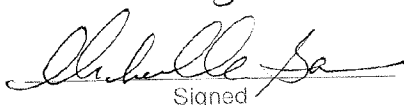
Inject mice w/ 100ul each i.p.

use 1cc syringe with 27 1/2 G needle

mice were not set up properly - we (Dan Scruggs + I) had to

redistribute mice to 5/cage before injection - mice were Stressed!
 injected mice 11:30am ✓ 1:30pm All OK ✓ 4:15pm all OK

Read and Understood By



Signed

Signed

Notebook Number: _____

Date: _____

9/6/02

✓ mice 8:00 Am

WT 5 - 3 mortalities

1 lethargic, shivering, unresponsive

1 Almost dead completely unresponsive, unable to walk euthanize w/CO₂
→ died before I could euthanize

WT 10 - 4 morts

1 sluggish but responsive

WT 50 3 mort

2 sluggish but responsive

WT 100 3 mort

2 sluggish but responsive have pilo erection

Dan2 500 All OK

100 all OK

50 all OK

10 all OK

PBS (-) control All OK

✓ mice 10:45 Am

WT 100 both mice are shivering, lethargic, very rapid breathing

✓ mice 2:45 pm

WT 100 breathing very rapidly unresponsive

WT 50 " "

WT 10 Very lethargic

WT 5 Sick but alert

Dan2 - all alert but not as alert as controls

✓ mice 6:20 pm

WT No change

Dan2 No change

PBS No change

Date: _____

10-4-02

Actinobacillus - plasmid + electroporation

- 1- Removed 40 μ l of actinobacillus competent cells from microfuge tube and added to a clean tube.
- 2- Added 1 μ l of plasmid DNA to the same tube.
- 3- Placed suspension in electroporation cuvette.
- 4- Set electroporator settings.
- 5- Electroporated and got a time constant = 5.26.
- 6- Placed cells suspension in 1 ml BHI_{NAD}⁺.
- 7- Incubated for 1 hour at 37°C.
- 8- Diluted culture to 10^{-2} .
- 9- Plated 100 μ l of 10^0 , 10^{-1} , 10^{-2} on BHI_{NAD}⁺ strept plates.
- 10- Incubated plates at 37°C overnight.

Results

10-5-02

Plate Count

10^0	33
10^{-1}	8
10^{-2}	0

Read and Understood By

Notebook Number: _____

Date: _____

10-29-02Mutation Rates

- 1- Diluted XLI Blue, XLI dam², Actin, and Actin dam² to 10^{-6} .
- 2- Plated in triplicate with 10 μ l of 10^{-5} and 10^{-6} .
- 3- Plated 100 μ l of overnight culture in triplicate on BHI, F. 100 μ l on BHI + AD⁺, F. 100 μ l on BHI + AD⁺.
- 4- Incubated at 37°C overnight.

Results

Plate Counts	10^{-5} TWTC	10^{-6} TWTC
XLI B		
XLI dam ²		
Actin		
Actin dam ²		

BHI, F

XLI	Bacterial lawn
XLI dam ²	Bacterial lawn
Actin	No growth
Actin dam ²	No growth

Note: Will have to do antibiotic titer with Actinobacillus and rifampin

10-29-02Fish Tissue (24 hour)

- 1- Pooled 3 fish from each tank.
- 2- Removed spleen and kidney.
- 3- Weighed, macerated, diluted to 10^{-2} .
- 4- Plated 15 μ l of 10^{-2} , 10^{-3} , 10^{-4} on EIM plates.
- 5- Incubated at 26°C for 48 hours.

Read and Understood By _____

1/30/03

1. Starter cultures of HB101 30.9% dand electroporation for plasmid prep.
2. Starter cultures of 93146 wt, R6, and EC 11229 for MK micro.

Conjugation with conjugation mutants

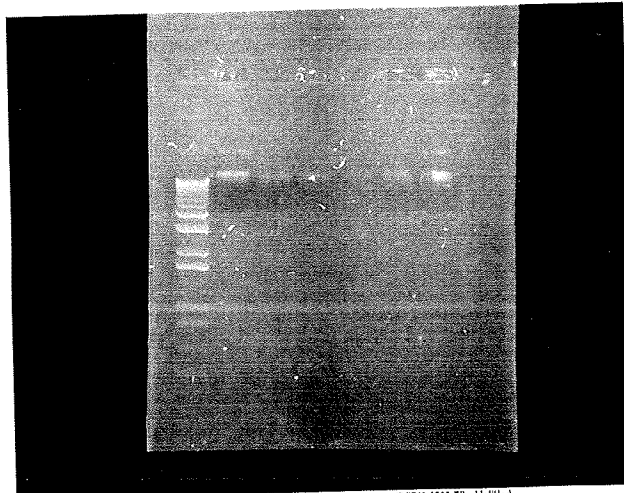
1. Pelleted bacteria in microtiter plate.
2. Resuspended in 100 μ l HBSS 0.9% with Ampicillin.
3. Removed 5 μ l from resuspended plate and added to 295 μ l HBSS μ mp.
4. Removed 5 μ l from the dilution plate and added to 35 μ l HBSS μ mp in white bottom assay plate.
5. Added an additional 40 μ l HBSS to central plate and 40 μ l normal serum to assay plate.
6. Incubated at RT for 1 hour.
7. Took readings on Dmax at Time 0, 30, and 60.
8. Analyzed data in Excel.
9. Highlighted mutants were from back (-80°C).

Results: See next page. (86)

1/30/03 HB101 30.9% dand plasmid prep.

1. Removed plasmid DNA with Qiagen kit.
2. Loaded 3 μ l on 7% agarose gel.
3. Ran at 100V for 1 hour.

Lane 1 - 1 kb ladder
 Lane 2 - prep 1
 Lane 3 - prep 2
 Lane 4 - prep 3
 Lane 5 - prep 4
 Lane 6 - prep 5
 Lane 7 - prep 6



Exp: 2.0 sec Bin: 1x1 Gain: 1.0 B:0 W:4871 G:0.55 N:0 Date: 30/12/2003 Time: 12:27:34 pm ID#513-2538 File: Untitled

B. J. P.
 Signed

Signed